## SEQUENCE LISTING

	<110> Cardio, Inc.	
5	<120> three dimensional tissue contruct	
	<130> CD009PCT	
	<150> JP2003-285476	
10	<151> 2003-08-01	
	<160> 20	
1 =	<170> Patentin version 3.1	
15	<210> 1	
	<211> 5956	
	<212> DNA	
20	⟨213⟩ Homo sapiens	
20	<400> 1	
	atgagttctg actcagaatt ggctgttttt ggggaggctg ctcctttcct ccgaaagtct	60
.25	gaaagggaac gcattgaggc ccagaatagg ccctttgatg ccaaaacatc tgtctttgtg	120
. <b>2</b> .9	goggagocca aagaatoott tgtcaaaggg accatocaga goagagaagg aggaaaagtg	180
	acggtgaaga ctgagggagg agcgactctg acagtgaagg atgatcaggt cttccccatg	240
30	aaccctccca aatatgacaa gatcgaggat atggccatga tgactcatct gcatgagcct	300
	gctgtgctgt acaacctcaa agaacgttat gcagcctgga tgatctacac ctattcaggt	360
35	ctcttctgtg tcactgtcaa cccctacaag tggctgcctg tgtataagcc cgaggtggtg	420
	acagoctaco gaggoaaaaa gogocaggag goocogococ acatottoto catototgac	480
	aacgcctatc agttcatgct gactgaccga gagaatcagt caatcctgat cactggagaa	540
40	totggtgcag ggaagactgt gaacaccaag cgtgtcatcc agtactttgc aacaattgca	600
	gttactggtg agaagaagaa ggaagaaatt acttctggca aaatacaggg gactctggaa	660
45	gatcaaatca tcagtgccaa cccctactg gaggcctttg gcaacgccaa gaccgtgagg	720
<del>4</del> 0	aatgacaact cctctcgctt tggtaaattc atcagaatcc actttggcac tactggaaaa	780
	ctggcatctg ctgatattga aacatatctg ctagagaagt ctagagttgt tttccagctt	RAN

	aaggctgaga	a gaagttatca	tattttta	cagattacat	cgaataagaa	accagaactt	900
5	attgaaatgo	ttotgattac	cacgaaccca	tatgattacc	catttgtcag	tcaaggggag	960
	atcagtgtgg	g ccagcatcga	tgatcaggaa	gaactgatgg	ccacagatag	tgctattgat	1020
	attttgggct	: ttactaatga	agaaaaggto	tccatttaca	agctcacggg	ggctgtgatg	1080
10	cattatggga	acctaaaatt	taagcaaaag	cagcgtgagg	agcaagcaga	gccagatggc	1140
	acagaagttg	ctgacaaggc	ggcctacctc	cagagtctga	actctgcaga	totgotoaaa	1200
15	gctctctgct	accccagggt	caaggtoggo	aatgagtatg	tcaccaaagg	ccagactgta	1260
	gaacaggtgt	ccaacgcagt	aggtgctctg	gccaaagccg	tctacgagaa	gatgttcctg	1320
	tggatggttg	cccgcatcaa	ccagcagctg	gacaccaagc	agcccaggca	gtacttcatc	1380
20	ggggtcttgg	acattgctgg	ttttgagatt	tttgatttca	acagcctgga	gcagctgtgc	1440
	atcaatttca	ccaatgagaa	actgcaacag	tttttcaacc	accacatgtt	cgtgctggag	1500
25	caggaggagt	acaagaagga	aggcatcgag	tggacgttca	togacttogg	gatggacctg	1560
	gctgcctgca	togagotoat	cgagaagcct	atgggcatct	tctccatcct	ggaagaggag	1620
	tgcatgttcc	ctaaggcaac	agacacctcc	ttcaagaaca	agctgtatga	ccagcacctg	1680
30	ggcaagtctg	ccaacttcca	gaagcccaag	gtggtcaaag	gcaaggccga	ggcccacttc	1740
	gctctgattc	actatgctgg	tgttgtggac	tacaacatta	ctggctggct	ggagaagaac	1800
35	aaggaccccc	tgaatgagac	cgtggttgga	ctgtaccaga	agtctgcaat	gaaaactcta	1860
	gctcagctct	tctctggggc	tcaaactgct	gaaggagagg	gagctggtgg	aggggccaag	1920
	aaaggtggta	agaagaaggg	ctcttcttc	cagacagtgt	ctgccctttt	cagagagaat	1980
40	ttgaacaagc	tgatgaccaa	cctcaggagt	acccatcctc	actttgtgag	gtgtatcatc	2040
	cccaatgaga	caaaaactcc	tggtgccatg	gagcatgagc	ttgtcctcca	ccagctgagg	2100
15	tgtaacggtg	tgctggaagg	catccgcatc	tgtaggaaag	gatttccaag	cagaatcctt	2160
	tatgcagact	tcaaacagag	atacaaggta	ttaaatgcaa	gtgcaatccc	tgaagggcaa	2220
	ttcattgata	gcaagaaggc	ctctgagaag	ctccttgcat	ccatcgacat ·	tgaccacacc .	2280

	cagtataaa	t ttgggcaca	c caaggtett	t ttcaaagct	g gtottotgg	g gctcctagag	2340
5	gagatgoga	g atgacaagc	t ggcccagct	g attacccgas	a cccaggcca	g gtgcagaggg	2400
	ttcttggca	a gagtggagt	a ccagaggat	g gtggagagaa	a gggaggcca	t cttctgtatc	2460
	cagtacaat	a tcagatoct	t catgaatgt	c aagcactggo	cctggatga	a actottotto	2520
10	aagatcaag	c ctctgttga	a gagtgcaga	a actgagaagg	g agatggccad	catgaaggaa	2580
	gaatttcaga	a aaattaaag	a cgaacttgc	c aagtcagagg	caaaaaggaa	a ggaactggaa	2640
15	gaaaagatgg	g tgacgctgt	t gaaagaaaa	a aatgacttgo	agctccaagt	tcaggctgaa	2700
	gccgaaggct	t tggctgatgo	o agaggaaagi	g tgtgaccagc	; taatcaaaac	caaaatccag	2760
	ctagaagcca	a aaatcaaaga	a ggtgactga	g agagotgagg	atgaggaaga	gatcaatgct	2820
20	gagotgacag	ccaagaagag	g gaaactggag	g gatgaatgtt	cagaactcaa	gaaagacatt	2880
	gatgaccttg	g agctgacact	ggccaaggti	t gagaaggaga	aacatgccac	agaaaacaag	2940
25	gtgaaaaacc	tcacagaaga	gatggcaggt	t ctggatgaaa	ccattgctaa	gctgaccaag	3000
	gagaagaagg	ctctccagga	ggcccaccag	cagaccctgg	atgacctgca	ggcagaggag	3060
	gacaaagtca	acaccctgac	caaagctaaa	atcaaacttg	aacaacaagt	ggatgatctt	3120
30	gaagggtcct	tggagcaaga	aaagaaactt	cgcatggacc	tagaaagggc	taagaggaaa	3180
	cttgagggtg	acttgaagtt	ggcccaagaa	tccataatgg	acattgaaaa	tgagaaacag	3240
35	caacttgatg	aaaagctcaa	aaagaaagag	tttgaaatca	gcaatctgca	aagcaagatt	3300
	gaagatgaac	aggcacttgg	cattcaattg	cagaagaaaa	ttaaagaatt	gcaagcccgc	3360
	attgaggagc	tggaggagga	aatcgaggcg	gagcgggcct	cccgggccaa	agcagagaag	3420
40	cagcgctctg	acctctcccg	ggagctggag	gagatcagcg	agaggotgga	agaagccggt	3480
	ggggccactt	cagoccagat	tgagatgaac	aagaagcggg	aggotgagtt	ccagaaaatg	3540
15	cgcagggaco	tggaggaggc	caccctacag	catgaagcca	cagcggccac	cctgaggaag	3600
-	aagcatgcag	atagtgtggc	cgagcttggg	gagcagattg	acaacctgca	gcgagtgaag	3660
	cagaagctgg	agaaggagaa	gagtgagatg	aagatggaga	ttgatgacct	tgctagtaat	3720

	gtagaaacg	g totocaaag	C Caagggaaa	c ctagagaaaa	a tgtgccggac	tctagaggac	3780
5	caactgagt	g aactgaaat	c aaaggaagaį	g gagcagcago	ggotgatoas	tgacctgact	3840
_	gcgcagagg	g ggcgcctgc	a gactgaatc	t ggtgagttti	cacgccagct	: tgatgaaaag	3900
	gaagetetg	g tgtctcagt	t atcaagaggo	aaacaagcct	ttactcaaca	gattgaagaa	3960
10	ttaaagagg	c aacttgaaga	a ggagataaaa	a gccaagaacg	g coctggogoa	tgccctgcag	4020
•	tettecege	c acgactgtga	a cctgctgcgg	gaacagtatg	aggaggagca	ggaatccaag	4080
15	googagotgo	agagagcact	gtccaaggco	aacaccgagg	ttgcccaatg	gaggaccaaa	4140
	tacgagacgg	g acgccatcca	gogoacagag	gagctggagg	aggccaagaa	gaagctggcc	4200
	cagoggotgo	aggcagctga	ı ggaacatgta	gaagctgtga	acgccaaatg	tgcttccctc	4260
20	gaaaagacga	agcagcggct	gcagaatgag	gtcgaggacc	tcatgottga	tgtggagagg	4320
	acaaatgccg	cctgtgccgc	ccttgacaaa	aagcaaagga	acttcgataa	gatoctggca	4380
25	gaatggaaac	agaaatgtga	ggaaacgcat	gctgagcttg	aggoctocca	gaaggaggcc	4440
	cgttcccttg	gcactgagct	gttcaagata	aagaatgcct	atgaggaatc	tttggatcag	4500
	ctagaaaccc	tgaagogaga	gaacaaaaac	ttacagcagg	agatttctga	cctcacggaa	4560
30	cagattgcag	aaggagggaa	acgtatccat	gaactggaga	aaataaagaa	acaagtggaa	4620
	caagaaaagt	gtgaacttca	ggctgcttta	gaagaagcag	aggcatctct	tgaacatgaa	4680
35	gagggaaaga	tcctgcgcat	ccagcttgag	ttgaaccaag	tcaagtctga	ggttgatagg	4740
		aaaaagatga					4800
	gagtccatgc	agagcacgct	ggatgctgag	atcaggagta	ggaatgatgc	cattaggctc	· 4860
40	aagaagaaga	tggagggaga	cctcaatgaa	atggaaatcc	agctgaacca	tgccaaccgc	4920
	atggctgctg	aggccctgag	gaactacagg	aacacccaag	gcatcctcaa	ggatacccag	4980
45	atccacctgg	atgatgctct	ccggagccag	gaggacctga	aggaacagct	ggccatggtg	5040
	gagogoagag	ccaacctgct	gcaggctgag	atcgaggagc	tgcgggccac	totggaacag	5100
	acagagagga	gcagaaaaat	cgcagaacag	gagctcctgg	atgccagtga ,	gogtgttcag	5160

	ctactgcaca cccagaacac cagcctgatc aacaccaaga agaagctgga gacagatatt	5220
5	toccaaatgo aaggagagat ggaggacatt otocaggaag coogcaatgo agaagaaaag	5280
Ū	gccaagaagg ccatcactga tgccgccatg atggctgagg agctgaagaa ggagcaggac	5340
	accagogoco acctggagog gatgaagaag aacatggago agacogtgaa ggatotgoag	5400
10	ctccgtctgg atgaggctga gcagctggcc ctgaagggtg ggaagaagca gatccagaaa	5460
	ctggaggcca gggtacggga gctggaagga gaggttgaga gtgagcaaaa gcgtaatgct	5520
15	gaggotgtca aaggtotgog caaacatgag aggogagtga aggaactcac ttaccagacg	5580
#.J	gaagaagata gaaagaatat totoaggott caagatttgg tagataaact toaggoaaaa	5640
	gtgaaatett ataagagaca agetgaggag getgaggaac aatecaacae caatetaget	5700
20	aaattoogoa agotooagoa tgagotggag gaggoogagg aacgggotga cattgotgag	5760
	tcccaggtga acaaactgcg ggtgaagagc cgggaggttc acacaaaagt cataagtgaa	5820
25	gagtgatcat gtcctgatgc catggaatga ctgaagacag gcacaaaatg tgacatcttt	5880
	ggtcatttcc ctctgtaatt attgtgtatt ctaccctgtt gcaaaggaaa taaagcatag	5940
	ggtagtttgc aaacaa	5956
30	(010)	
	<210> 2	
	⟨212⟩ PRT	
	<213> Homo sapiens	
35		
	<b>&lt;400&gt;</b> 2	
	Met Ser Ser Asp Ser Glu Leu Ala Val Phe Gly Glu Ala Ala Pro Phe	
10	1 5 10 15	
	Leu Arg Lys Ser Glu Arg Glu Arg 11e Glu Ala Gln Asn Arg Pro Phe 20 25 30	
15		

Asp Ala Lys Thr Ser Val Phe Val Ala Glu Pro Lys Glu Ser Phe Val

45

	Ly	s GI; 50	y Thi	- 1.16	e Gin	Ser	55 55	g Glu	Gly	Gly	/ Lys	60	Thr	· Val	Lys	Thr
5	G1u 65	ı Gl	y Gly	/ Ala	1 Thr	Leu 70	1 Thr	·Val	Lys	Asp	Asp 75	Gin	Val	Phe	Pro	Met 80
10	Asr	n Pro	Pro	Lys	Tyr 85	Asp	Lys	lle	Glu	Asp 90	Met	Ala	Met	Met	Thr 95	His
15	Leu	His	s Glu	Pro 100		Val	Leu	Tyr	Asn 105		Lys	Glu	Arg	Tyr 110		Ala
20	Trp	Met	: lle 115	Tyr	Thr	Tyr	Ser	Gly 120		Phe	Cys	Val	Thr 125		Asn	Pro
	Tyr	Lys 130	Trp	Leu	Pro	Val	Tyr 135	Lys	Pro	Glu	Val	Va I 140	Thr	Ala	Tyr	Arg
25	Gly 145	Lys	Lys	Arg	GIn	Glu 150	Ala	Pro	Pro	His	lle 155	Phe	Ser	lie	Ser	Asp 160
30	Asn	Ala	Tyr	Gln	Phe 165	Met	Leu	Thr	Asp	<b>A</b> rg 170	Glu	Asn	GIn	Ser	lle 175	Leu
35	lle	Thr	Gly	GIu 180	Ser	Gly	Ala	Gly	Lys 185	Thr	Val	Asn	Thr	Lys 190	Arg	Val
40	lle	Gin	Tyr 195	Phe	Ala	Thr	lle	Ala 200	Val	Thr	Gly	Glu	Lys 205	Lys	Lys	Glu
	Glu	lle 210	Thr	Ser	Gly	Lys	lle 215	Gln	Gly	Thr	Leu	Glu 220	Asp	Gin	lle	He
45	Ser 225	Ala	Asn	Pro	Leu	Leu 230	Glu	Ala	Phe		Asn 235	Ala	Lys	Thr	Val	Arg 240

	Asn	ı Asp	) Ası	n Ser	245		Phe	Gly	Lys	250		Arg	; lle	His	Phe 255	
5	Thr	Thr	· GI)	/ Lys 260		ı Ala	Ser	· Ala	Asp 265		Glu	Thr	Tyr	Leu 270		Glu
10	Lys	Ser	Arg 275		Val	Phe	Gin	Leu 280		Ala	Glu	Arg	Ser 285		His	lle
15	Phe	Tyr 290		ılle	Thr	Ser	Asn 295	Lys	Lys	Pro	Glu	Leu 300		Glu	Met	Leu
20	Leu 305		Thr	Thr	· Asn	Pro 310	Tyr	Asp	Tyr	Pro	Phe 315	Val	Ser	GIn	Gly	Glu 320
	He	Ser	Val	Ala	Ser 325	lle	Asp	Asp	Gln	Glu 330	Glu	Leu	Met	Ala	Thr 335	Asp
25	Ser	Ala	He	Asp 340		Leu	Gly	Phe	Thr 345	Asn	Glu	Glu	Lys	Va I 350	Ser	lle
30	Tyr	Lys	Leu 355	Thr	Gly	Ala	Val	Met 360	His	Tyr	Gly	Asn	Leu 365	Lys	Phe	Lys
35	Gin	Lys 370	Gin	Arg	Glu	Glu	Gin 375	Ala	Glu	Pro	Asp	Gly 380	Thr	Glu	Val	Ala
10	Asp 385	Lys	Ala	Ala	Tyr	Leu 390	Gin	Ser	Leu	Asn	Ser 395	Ala	Asp	Leu	Leu	Lys 400
	Ala	Leu	Cys	Tyr	Pro 405	Arg	Val	Lys	Val	Gly 410	Asn	Glu	Tyr	Val	Thr 415	Lys
15	Gly	Gin	Thr	Va I 420	Glu	Gln	Va!		Asn 425	Ala	Val	Gly	Ala	Leu 430	Ala	Lys

	Ala	Val	1yr 435		ı Lys	Met	Phe	Leu 440		Met	Val	Ala	Arg 445		Asn	Gin
5	Gin	Leu 450	Asp	Thr	· Lys	GIn	Pro 455		GIn	Tyr	Phe	11e 460	Gly	Val	Leu	Asp
10	1 (e 465		Gly	Phe	Glu	11e 470		Asp	Phe	Asn	Ser 475		Glu	Gin	Leu	Cys 480
15	He	Asn	Phe	Thr	Asn 485		Lys	Leu	Gin	GIn 490	Phe	Phe	Asn	His	His 495	
20	Phe	Val	Leu	Glu 500		Glu	Glu	Tyr	Lys 505	Lys	Glu	Gly	lle	Glu 510	Trp	Thr
	Phe	lle	Asp 515	Phe	Gly	Met	Asp	Leu 520	Ala	Ala	Суз	lle	Glu 525	Leu	lle	Glu
25	Lys	Pro 530	Met	Gly	lle	Phe	Ser 535	lle	Leu	Glu	Glu	Glu 540	Cys	Met	Phe	Pro
30	Lys 545	Ala	Thr	Asp	Thr	Ser 550	Phe	Lys	Asn	Lys	Leu 555	Tyr	Asp	GIn	His	Leu 560
35	Gly	Lys	Ser	Ala	Asn 565	Phe	GIn	Lys	Pro	Lys 570	Val	Val	Lys	Gly	Lys 575	Ala
.40	Glu	Ala	His	Phe 580	Ala	Leu	He	His	Tyr 585	Ala	Gly	Val	Val	Asp 590	Tyr	Asn
	lle	Thr	Gly 595	Trp	Leu	Glu	Lys	Asn 600	Lys	Asp	Pro	Leu	Asn 605	Glu	Thr	Val
45	Val	Gly 610	Leu	Tyr	Gin	Lys	Ser 615	Ala	Met	Lys	Thr	Leu 620	Ala	Gln	Leu	Phe

	Ser 625	. G17	/ Ala	ı Gir	1 Thr	630		ı Gly	Glu	ı Gly	Ala 635		Gly	Gly	Ala	Lys 640
5	Lys	Gly	gly	Lys	Lys 645		Gly	' Ser	Ser	Phe 650		Thr	Val	Ser	Ala 655	
10	Phe	Arg	; Glu	Asn 660		Asn	Lys	Leu	Met 665		Asn	Leu	Arg	Ser 670		His
15	Pro	His	Phe 675		Arg	Cys	He	11e 680		Asn	Glu	Thr	Lys 685	Thr	Pro	Gly
20	Ala	Met 690	Glu	His	Glu	Leu	Va I 695		His	Gin	Leu	Arg 700		Asn	Gly	Val
	Leu 705		Gly	He	Arg	l le 710	Cys	Arg	Lys	Gly	Phe 715	Pro	Ser	Arg	l le	Leu 720
25	Tyr	Ala ·	Asp	Phe	Lys 725	Gln	Arg	Tyr	Lys	Va I 730	Leu	Asn	Ala	Ser	Ala 735	lie
30	Pro	Glu	Gly	GIn 740		lle	Asp	Ser	Lys 745		Ala	Ser	Glu	Lys 750	Leu	Leu
35	Ala	Ser	lle 755	Asp	He	Asp	His	Thr 760	Gln	Tyr	Lys	Phe	Gly 765	His	Thr	Lys
40	Val	Phe 770	Phe	Lys	Ala	Gly	Leu 775	Leu	Gly	Leu	Leu	Glu 780	Glu	Met	Arg	Asp
	Asp 785	Lys	Leu	Ala	GIn	Leu 790	He	Thr	Arg	Thr	GIn 795	Ala	Arg	Cys	Arg	Gly 800
45	Phe	Leu	Ala	Arg	Va I 805	Glu	Tyr	Gln	Arg	Met 810	Val	Glu	Arg	Arg	Glu 815	Ala

	lle	Phe	Сув	11e 820		Tyr	Asn	lle	Arg 825	Ser	Phe	Met	Asn	Va I 830	Lys	His
5	Trp	Pro	Trp 835	Met	Lys	Leu	Phe	Phe 840	Lys	lle	Lys	Pro	Leu 845		Lys	Ser
10	Ala	GIu 850		Glu	Lys	Glu	Met 855	Ala	Thr	Met	Lys	<b>G</b> lu 860	Glu	Phe	Gln	Lys
15	11e 865		Asp	Glu	Leu	Ala 870	Lys	Ser	Glu	Ala	Lys 875	Arg	Lys	Glu	Leu	Glu 880
20	Glu	Lys	Met	Va I	Thr 885	Leu	Leu	Lys	Glu	Lys 890	Asn	Asp	Leu	Gin	Leu 895	
	Val	Gln	Ala	G1u 900	Ala	Glu	Gly	Leu	Ala 905	Asp	Ala	Glu	Glu	Arg 910	Cys	Asp
25	GIn	Leu	lle 915	Lys	Thr	Lys	lle	GIn 920	Leu	Glu	Ala	Lys	lle 925	Lys	Glu	Val
30	Thr	Glu 930	Arg	Ala	Glu	Asp	Glu 935	Glu	Glu	He	Asn	Ala 940	Glu	Leu	Thr	Ala
35	Lys 945	Lys	Arg	Lys	Leu	Glu 950	Asp	Glu	Cys	Ser	Glu 955	Leu	Lys	Lys	Asp	lle 960
40	Asp	Asp	Leu	Glu	Leu 965	Thr	Leu	Ala	Lys	Va I 970	Glu	Lys	Glu	Lys	His 975	Ala
	Thr	Glu	Asn	Lys 980	Val	Lys	Asn		Thr 985	Glu	Glu	Met	Ala	Gly 990	Leu	Asp
45	Glu	Thr	lle 995	Ala	Lys	Leu		Lys 1000		Lys	Lys	: Ala	100		n Gl	u Ala

	His Gir 101	ı Gir O	1 Thr	· Leu	ı Asp	Asp 101		u Glr	n Ala	Gľu	1020		Lys	s Va
5	Asn Thr 102	Leu 5	ı Thr	Lys	Ala	Lys 103(		e Lys	s Leu	· Glu	GIn 1035		Val	Ast
10	Asp Leu 104	Glu O	ı Gly	Ser	Leu	Glu 1045	Glr	n Glu	ı Lys	Lys	Leu 1050		Met	: Asp
15	Leu Glu 105	Arg 5	: Ala	Lys	Arg	Lys 1060		ı Glu	ı Gly	Asp	Leu 1065		Leu	Ala
20	Gin Giu 1070	Ser )	lle	Met	Asp	lle 1075		Asn	Glu	Lys	01n 1080	GIn	Leu	Asp
٠	Glu Lys 1085	Leu	Lys	Lys	Lys	Glu 1090	Phe	Glu	lle	Ser	Asn 1095	Leu	Gin	Ser
25	Lys IIe 1100	Glu	Asp	Glu	GIn	Ala 1105	Leu	Gly	lle	GIn	Leu 1110	Gln	Lys	Lys
30	lle Lys 1115	Glu	Leu	Gln	Ala	Arg 1120	ile	Glu	Glu	Leu	Glu 1125	Glu	Glu	lle
35	Glu Ala 1130	Glu	Arg	Ala		Arg 1135	Ala	Lys	Ala		Lys 1140	Gln	Arg	Ser
40	Asp Leu 1145	Ser	Arg	Glu i	Leu	Glu 1150	Glu	He	Ser		Arg 1155	Leu	Glu	Glu
•	Ala Giy 1160	Gly	Ala '	Thr S	Ser .	Ala 1165	Gin	lle	Glu		Asn 1170	Lys I	Lys	Arg
45	Glu Ala 1175	Glu l	Phe (	3in (	_ys l	Met 1180	Arg	Arg .	Asp		91u ( 1185	Glu .	Ala	Thr

	Le	u Gin 119	Hi: O	s Glu	J Ala	1 Thr	Ala 1195		1 Thr	· Leu	Arg	Lys 1200		His	Ala
5	As	p Ser 120	<b>Va</b> ↓ 5	l Ala	Glu	l Leu	1210	Glu )	ı Gin	lle	Asp	Asn 1215		Gin	Arg
10	Va	l Lys 1220		n Lys	Leu	Glu	Lys 1225		Lys	Ser	Glu	Met 1230		Met	Glu
15	He	e Asp 1238	Asp 5	Leu	Ala	Ser	Asn 1240		Glu	Thr	Val	Ser 1245		Ala	Lys
20	GI	/ Asn 1250		Glu	Lys	Met	Cys 1255		Thr	Leu	Glu	Asp 1260	GIn	Leu	Ser
	Glu	1 Leu 1265	Lys	Ser	Lys	Glu	Glu 1270		GIn	GIn	Arg	Leu 1275	He	Asn	Asp
25	Leu	Thr 1280		Gln	Arg	Gly	Arg 1285		GIn	Thr	Glu	Ser 1290	Gly	Glu	Phe
30	Ser	Arg 1295		Leu	Asp	Glu	Lys 1300	Glu	Ala	Leu	Val	Ser 1305	GIn	Leu	Ser
35	Arg	Gly 1310	Lys	GIn	Ala	Phe	Thr 1315	<b>G</b> In	GIn	He	Glu	Glu 1320	Leu	Lys	Arg
40	GIn	Leu 1325	Glu	Glu	Glu	lle	Lys 1330	Ala	Lys	Asn		Leu 1335	Ala	His	Ala
	Leu	GIn 1340	Ser	Ser	Arg		Asp 1345	Cys	Asp	Leu		Arg 1350	Glu	Gin	Tyr
45	Glu	Glu 1355	Glu	Gin	Glu		Lys 1360	Ala	Glu	Leu		Arg 1365	Alal	Leu :	Ser

	Lys	1370		Thr	· Glu	ı Val	Ala 1375		Trp	Arg	Thr	Lys 1380		Glu	Thr
5	Asp	Ala 1385		Gin	Arg	: Thr	Glu 1390		Leu	Glu	Glu	Ala 1395		Lys	Lys
10	Leu	Ala. 1400		Arg	Leu	Gln	Ala 1405		Glu	Glu	His	Val 1410		Ala	Val
15	Asn	Ala 1415		Cys	Ala	Ser	Leu 1420		Lys	Thr	Lys	GIn 1425	Arg	Leu	GIn
20	Asn	Glu 1430		Glu	Asp	Leu	Met 1435		Asp	Val	Glu	Arg 1440	Thr	Asn	Ala
	Ala	Cys` 1445		Ala	Leu	Asp	Lys 1450		Gln	Arg	Asn	Phe 1455	Asp	Lys	He
25	Leu	Ala 1460		Trp	Lys	61n	Lys 1465		Glu	Glu	Thr	His 1470	Ala	Glu	Leu
30	Glu	Ala 1475		Gin	Lys	Glu	Ala 1480	Arg	Ser	Leu	Gly	Thr 1485	Glu	Leu	Phe
35	Lys	lle 1490		Asn	Ala	Tyr	Glu 1495	Glu	Ser	Leu	Asp	Gln 1500	Leu	Glu	Thr
40	Leu	Lys 1505	Arg	Glu	Asn	Lys	Asn 1510	Leu	GIn	GIn	Glu	l le 1515	Ser	Asp	Leu
	Thr	Glu 1520	GIn	He	Ala	Glu	Gly 1525	Gly	Lys	Arg	lle	His 1530	Glu	Leu	Glu
45	Lys	lle 1535	Lys	Lys	Gin	Val <sub>.</sub>	Glu 1540	GIn	Glu	Lys	Cys	Glu 1545	Leu	Gin	Ala

	Ala	Leu 1550		Glu	Ala	Glu	Ala 1555		Leu	Glu	His	Glu 1560	Glu	Gly	Lys
5	lle	Leu 1565		lle	GIn	Leu	Glu 1570		Asn	GIn	Val	Lys 1575	Ser	Glu	Val
10	Asp	Arg 1580		lle	Ala	Glu	Lys 1585		Glu	Glu	He	Asp 1590	Gin	Leu	Lys
15	Arg	Asn 1595	His	He	Arg	ile	Val 1600		Ser	Met	Gin	Ser 1605	Thr	Leu	Asp
20	Ala	Glu 1610	Ile	Arg	Ser	Arg	Asn 1615		Ala	lle	Arg	Leu 1620	Lys	Lys	Lys
	Met	Glu 1625	Gly	Asp	Leu	Asn	Glu 1630	Met	Glu	He	GIn	Leu 1635	Asn	His	Ala
25	Asn	Arg 1640	Met	Ala	Ala	Glu	Ala 1645	Leu	Arg	Asn	Tyr	Arg 1650	Asn	Thr	GIn
30	Gly	lle 1655	Leu	Lys	Asp	Thr	GIn 1660	lle	His	Leu	Asp	Asp 1665	Ala	Leu	Arg
35	Ser	GIn 1670	Glu	Asp	Leu	Lys	Glu 1675	Gin	Leu	Ala	Met	Val 1680	Glu	Arg	Arg
40	Ala	Asn 1685	Leu	Leu	Gln	Ala	Glu 1690	lle	Glu	Glu	Leu	Arg 1695	Ala	Thr	Leu
	Glu	Gln 1700	Thr	Glu	Arg	Ser	Arg 1705	Lys	He	Ala	Glu	GIn 1710	Glu	Leu	Leu
45	Asp	Ala 1715	Ser	Glu	Arg	Vai	GIn 1720	Leu	Leu	His	Thr	GIn 1725	Asn	Thr	Ser

	Leu	1730		1 Thr	Lys	: Lys	Lys 1735		Glu	Thr	Asp	11e 1740		Gin	Met
<sup>.</sup> 5	GIn	1745		Met	Glu	Asp	lle 1750		GIn	Glu	Ala	Arg 1755		Ala	Glu
10	Glu	Lys 1760		Lys	Lys	Ala	lle 1765		Asp	Ala	Ala	Met 1770		Ala	Glu
15		Leu 1775		Lys	Glu	GIn	Asp 1780		Ser	Ala	His	Leu 1785		Arg	Met
20	Lys	Lys 1790		Met	Glu	Gln	Thr 1795		Lys	Asp	Leu	GIn 1800	Leu	Arg	Leu
	Asp	GIu 1805		Glu	GIn	Leu	Ala 1810		Lys	Gly	Gly	Lys 1815	Lys	Gln	lle
25	GIn	Lys 1820		Glu	Ala	Arg	Va I 1825		Glu	Leu	Glu	Gly 1830	Glu	Val	Glu
30	Ser	Glu 1835		Lys	Arg	Asn	Ala 1840	Glu	Ala	Val	Lys	Gíy 1845	Leu	Arg	Lys
35	His	Glu 1850	Arg	Arg	Val	Lys	Glu 1855	Leu	Thr	Tyr	GIn	Thr 1860	Glu	<b>Glu</b>	Asp
40	Arg	Lys 1865	Asn	He	Leu	Arg	Leu 1870	GIn	Asp	Leu	Val	Asp 1875	Lys	Leu	GIn
	Ala	Lys 1880	Val	Lys	Ser	Tyr	Lys 1885	Arg	GIn	Ala	Glu	Glu 1890	Ala	Glu	Glu
15	GIn	Ser 1895	Asn	Thr	Asn	Leu	Ala 1900	Lys	Phe	Arg	Lys	Leu 1905	Gin	His	Glu

Glu 1910	Glu	Ala	Glu	Glu	Arg 1915	Asp	He	Ala	Glu 1920	Gin	Val

Asn Lys Leu Arg Val Lys Ser Arg Glu Val His Thr Lys Val He 1925 1930 1935

10 Ser Glu Glu 1940

**<400>** 3

20 atcottocto aaaattottg aagtagttgt ctgctttgag cctgccacct tottcatctg 60 ataatacaag aggtatacct agtccagcac tgccatcaat aacctgcagc catgagttct 120 gactotgaga tggccatttt tggggaggot gotoctttoc tocgaaagto tgaaaaggag 180 25 cgaattgaag ctcagaacaa gccttttgat gccaagacat cagtctttgt ggtggaccct 240 aaggagtoot acgtgaaagc aatagtgcag agcagggaag gggggaaggt gacagccaag 300 30 accgaagetg gagetactgt aactgtgaaa gaagaccaag tetteteeat gaacceteee 360 420 tataacctca aagagcgtta cgcagcctgg atgatctaca cctactcggg cctcttctgt 480 35 gtcaccgtca acccctacaa gtggctgccg gtgtacaacc ctgaggtggt gacagcctac 540 cgaggcaaaa agcgccagga ggccccaccc catatettet ccatetetga caatgcetat 600 40 cagttcatgc taactgatcg tgaaaaccag tcaatcttga ttactggaga atctggtgca 660 gggaagactg tgaacacgaa gogtgtcatc cagtactttg caacaattgc agttactgga 720 gagaagaaaa aagaggaacc tgcctctggc aaaatgcagg ggacccttga agatcaaatc 780 45 atcagtgcta accccctact ggaagccttc ggcaatgcca agaccgtgag gaatgacaac 840 toctotogot tiggiaaatt catcaggato cattitggig ccacaggcaa aciggottot 900

,	gcagatatt	g aaacatatc	t gotagagaag	tcccgagtta	cttttcagc	t aaaggotgaa	960
5	agaagctac	c acatatttt	a tcaaatcctg	tccaataaga	aaccagagc	cattgaaatg	1020
-	cttctgatc	a ccaccaacco	atatgactto	gcatttgtca	gccaagggg	a aattactgtg	1080
	cccagcatt	g atgaccagge	a agagctgatg	gccacagata	gtgctgtggg	catcctgggt	1140
10	ttcactgct	g atgaaaaggt	ggccatttac	aagctcactg	gagccgtgat	gcattatggg	1200
	aacatgaaa <sup>.</sup>	t tcaagcaaaa	gcaaagggaa	gagcaggcag	agccagatgg	cacggaagtt	1260
15	gctgacaaa	g ctgcttatct	gacaagtctg	aactctgctg	acctgctcaa	atctctctgc	1320
	tatcccaga	g tcaaggtcgg	caatgagttc	gtaaccaaag	gccagactgt	gcagcaggtg	1380
	tacaacgcag	g tgggtgctct	ggccaaagcc	atctacgaga	agatgttcct	gtggatggtc	1440
20	acccgcatca	a accagcaget	ggacaccaag	cagcccaggc	agtacttcat	cggggtcttg	1500
	gacattgctg	gctttgagat	ctttgatttc	aacagcctgg	agcagctgtg	catcaacttc	1560
25	accaacgaga	aactgcaaca	gtttttcaac	caccacatgt	togtgotgga	gcaggaagag	1620
		aaggcatcga					1680
		tcgagaagcc					1740
30		cagacacctc					1800
		agaagcccaa			•		1860
35		gcaccgtgga					1920
		ctgtggtggg					1980
		cacaaactgc					2040
40	•	ctttccagac					2100
	accaacttga	ggagcactca	ccccacttt	gtgcggtgca	tcatccccaa	tgaaactaaa	2160
45	actcctggtg	ccatggagca	tgagcttgtc	ctgcatcagc	tgaggtgtaa	cggtgtgctg	2220
	gaaggcatcc	gcatctgcag	gaaaggcttc	ccaagcagaa	tcctttatgc	agacttcaaa	2280
	cagagataca	aggttctaaa	tgcgagtgct	atcccagagg .	gtcagttcat	tgacagcaag	2340

	aaggottote	g agaaacttct	agggtotatt	gaaattgaco	acacocagta	caaattcggt	2400
5	cataccaage	ttttcttcae	agctggcctg	ctgggaactc	tagaagaaat	gogagatgaa	2460
J	aagotagoto	aactcatcac	gogoactcaa	gccatatgca	gggggttcct	gatgagagtg	2520
	gagttcagae	agatgatgga	gaggagagag	tccatcttct	gcattcagta	caacatccgt	2580
10	gctttcatga	atgtgaagca	ctggccctgg	atgaagctgt	atttcaagat	caagcccctc	2640
	ctcaagagtg	cagagacaga	gaaggagatg	gccaacatga	aggaagaatt	tgagaaaacc	2700
15	aaagaagago	tggctaagac	agaggcaaaa	aggaaagaac	tagaagaaaa	gatggtgacg	2760
	ctaatgcaag	agaaaaatga	cttacaactc	caagttcaag	ctgaagcaga	tgccttggct	2820
	gatgcagagg	aaagatgtga	tcagttgatt	ававссавав	tccaacttga	ggccaaaatc	2880
20	aaagaggtaa	ctgaaagagc	tgaggatgag	gaagagatca	atgotgagot	gacagccaag	2940
	aagaggaaac	tggaggatga	atgttcagag	ctcaagaaag	acattgatga	ccttgagctg	3000
25	acactggcca	aggttgagaa	ggagaaacat	gccacagaga	acaaggtgaa	aaacctcaca	3060
	gaagagatgg	caggtotgga	tgaaaccatt	gctaagctga	ccaaggagaa	gaaggctctc	3120
	caggaggccc	accagcagac	cctggatgac	ctgcagatgg	aggaggacaa	agtcaacacc	3180
30	ctgaccaaag	ctaaaaccaa	gctagaacag	caagtggacg	atcttgaagg	atctctggaa	3240
	caagaaaaga	aactttgcat	ggacttagaa	agagccaaga	gaaaactgga	gggtgaccta	3300
35	aaattggccc	aagaatccac	aatggataca	gaaaatgaca	aacagcaact	taatgagaaa	3360
	ctcaaaaaga	aagagtttga	aatgagcaat	ctgcaaggca	agattgaaga	tgaacaagcc	3420
	cttgcaatgc	agctacaaaa	gaagatcaaa	gaattacagg	cccgcattga	ggagctggag	3480
40	gaggaaatcg	aggcagagcg	ggcctcccgg	gccaaagcag	aaaagcagcg	ctctgacctc	3540
	tcccgggagc	tggaggagat	cagtgagagg	ctggaagaag	ccggtggggc	cacttcagcc	3600
<b>4</b> 5 .	cagattgagt	tgaacaagaa	gcgggaggct	gagttccaga.	aaatgcgcag	ggacctggaa	3660
	gagtccaccc	tgcagcacga	agocacggca	gctgctcttc	ggaagaagca	cgcagatagt	3720
	gtggctgagc	ttgggaagca	gatogacago	cttcagcggg	tcaagcagaa	gctggagaag	3780

	gaaaagagt	g agctgaaga	t ggagatcaa	t gaccttgcta	a gtaacatgga	gactgtctcc	3840
5	aaagccaag	g caaactttg	a gaaaatgtg	c cgcaccctag	g aggaccagct	tagtgaaata	3900
Ū	aaaacaaag	g aagaagagc	a gcaacgott	a ataaatgagi	tgtcagccca	gaaggcacgt	3960
	ttacacacag	g aatcaggtga	a gttttcacga	a cagctagate	g aaaaagatgo	tatggtttct	4020
10	cagctatcc	gaggcaaaca	a agcatttaca	a caacagatte	, aagaattaaa	gaggcagcta	4080
	gaagaggaga	a ctaaggccaa	a gagcactote	g gcccatgcco	tgcagtcagc	ccgccatgac	4140
15	tgtgacctgo	tgcgggaaca	a gtatgaggag	gagcaggaag	ccaaggotga	gctgcagagg .	4200
	ggaatgtcca	aggccaacag	tgaggttgco	cagtggagga	ccaagtacga	gacggacgcc	4260
	atccagcgca	cagaggagci	: ggaggaggco	aagaagaagc	tagcccagcg	tctgcaggat	4320
20	gcagaagaac	atgtagaago	; tgtgaattco	aaatgtgott	ctcttgaaaa	gacaaagcag	4380
	aggctacaga	atgaagtaga	ggacctcatg	attgatgtgg	aacgatctaa	tgctgcctgc	4440
25	atageteteg	ataagaagca	aagaaacttt	gacaaggttc	tggcagaatg	gaaacagaag	4500
	tatgaggaaa	ctcaggctga	acttgaggco	tcccagaagg	agtogogtto	tctcagcact	4560
					atcatcttga		4620
30					cagagcaaat	•	4680
					ttgatcatga		4740
35					atgaagaagg		4800
	•				accgaaaaat		4860
					ttgtggagtc		4920
40					ggatcaagaa		4980
					accgccaggc		5040
15					ctcagctaca	•	5100
					tggttgagcg		5160
	ctgatgcagg	ctgaagttga	agagotoagg	gcatccctgg	aacggactga (	gagaggcagg	5220

actgatgctg ccatgatggc tgaggagctg aagaaggaac aggacaccag cgcccacctg 54  10 gagcggatga agaagaacat ggagcagacc gtgaaggatc tgcagctccg tctgggtgag 55  gctgagcagc tggcgctgaa gggtgggaag aagcagatcc agaaactgga ggccagggtg 55  agaagacttg aaagtgaggt ggaaagtgaa cagaagcaca atgttgaggc tgtcaagggt 56  cttcgcaaac atgagagaag agtgaaggaa ctcacttacc agactgagga ggaccgcaag 57	<b>4</b> 0
gagatgagg acatcgtcca ggaagcccgc aatgcagagg agaaggccaa gaaggccatc 54 actgatgctg ccatgatggc tgaggagctg aagaaggaac aggacaccag cgcccacctg 54 10 gagcggatga agaagaacat ggagcagacc gtgaaggatc tgcagctccg tctgggtgag 55 gctgagcagc tggcgctgaa gggtgggaag aagcagatcc agaaactgga ggccagggtg 55 agaagacttg aaagtgaggt ggaaagtgaa cagaagcaca atgttgaggc tgtcaagggt 56 cttcgcaaac atgagagaag agtgaaggaa ctcacttacc agactgagga ggaccgcaag 57	-10
getgageage tggegetgaa gggtgggaag aageagate tgeageteeg tetgggtgag 55 getgageage tggegetgaa gggtgggaag aageagatee agaaactgga ggeeagggtg 55 agaagagettg aaagtgaggt ggaaagtgaa cagaageaca atgttgagge tgteaagggt 56 cttegeaaac atgagagaag agtgaaggaa eteaettace agactgagga ggacegeaag 57	00
gctgagcagc tggcgctgaa gggtgggaag aagcagatcc agaaactgga ggccagggtg 55 agagagcttg aaagtgaggt ggaaagtgaa cagaagcaca atgttgaggc tgtcaagggt 56 15 cttcgcaaac atgagagaag agtgaaggaa ctcacttacc agactgagga ggaccgcaag 57	60
agagagcttg aaagtgaggt ggaaagtgaa cagaagcaca atgttgaggc tgtcaagggt 56  15  cttcgcaaac atgagagaag agtgaaggaa ctcacttacc agactgagga ggaccgcaag 57	20
15 cttcgcaaac atgagagaag agtgaaggaa ctcacttacc agactgagga ggaccgcaag 57	80
cttcgcaaac atgagagaag agtgaaggaa ctcacttacc agactgagga ggaccgcaag 57	40
	00
aatattotoa ggotgoagga ottggtggac aaattgcaaa ocaaagtoaa agottacaag 57	60
20 agacaagctg aagaggctga ggaacaatcc aatgtcaacc ttgccaagtt ccgcaagctc 58	20
cagcacgagc tggaggaggc cgaggaacgg gctgacattg ctgagtccca agtcaacaag 58	80
ctgagagtga agagtcggga ggttcacaca aaagtcataa gtgaagagta attcattcta 59	40
atgaaagaaa atgtgaccaa agaaatgcac gaaatgtgaa gttctttgtc actgtcctgt 60	00
atatcaagga aataaa 60	16
30 <b>&lt;210&gt; 4</b>	
<210> 4 <211> 1939	
<212> PRT	
<213> Homo sapiens	
35 <b>&lt;400&gt; 4</b>	
Met Ser Ser Asp Ser Glu Met Ala IIe Phe Gly Glu Ala Ala Pro Phe	
1 5 10 15 40	
Leu Arg Lys Ser Glu Lys Glu Arg lie Glu Ala Gln Asn Lys Pro Phe 20 25 30	
<b>4</b> 5	

Asp Ala Lys Thr Ser Val Phe Val Val Asp Pro Lys Glu Ser Tyr Val

45

40

	Lys	50	l IIe	Val	Gin	Ser	Arg 55	Glu	Gly	Gly	Lys	Va I 60	Thr	Ala	Lys	Thr
5	G1 u 65	ı Ala	Gly	Ala	Thr	Va I 70	Thr	Val	Lys	Glu	Asp 75	GIn	Val	Phe	Ser	Met 80
10	Asn	Pro	Pro	Lys	Tyr 85	Asp	Lys	lle	Glu	Asp 90	Met	Ala	Met	Met	Thr 95	His
15	Leu	His	Glu	Pro 100		Val	Leu	Tyr	Asn 105	Leu	Lys	Glu	Arg	Tyr 110	Ala	Ala
20	Trp	Met	11e 115	Tyr	Thr	Tyr	Ser	Gly 120	Leu	Phe	Cys	Val	Thr 125	Val	Asn	Pro
	Tyr	Lys 130		Leu	Pro	Val	Tyr 135	Asn	Pro	Glu	Val	Va I 140	Thr	Ala	Tyr	Arg
25	Gly 145		Lys	Arg	Gln	Glu 150	Ala	Pro	Pro	His	l le 155	Phe	Ser	He	Ser	<b>Asp</b> 160
30	Asn	Ala	Tyr	Gin	Phe 165	Met	Leu	Thr	Asp	Arg 170	Glu	Asn	Gin	Ser	l le 175	Leu
35	lle	Thr	Gly	<b>G</b> lu 180	Ser	Giy	Ala	Gly	Lys 185	Thr	Val	Asn	Thr	Lys 190	Arg	Val
40	He	Gin	Tyr 195	Phe	Ala	Thr	He	Ala 200	Val	Thr	Gly	Glu	Lys 205	Lys	Lys	Glu
	Glu	Pro 210	Ala	Ser	Gly	Lys	Met 215	GIn	Gly	Thr	Leu	Glu 220	Asp	GIn	lle	He
45	Ser 225	Ala	Asn	Pro	Leu	Leu 230	Glu	Ala	Phe	Gly	Asn 235	Ala	Lys	Thr	Val	Arg 240

·	AS	II AS	SP AS	sn 90	24!		g Pne	e Gij	y Lys	250		Are	g ile	His	255	•
5	AI.	a Th	ır GI	y Ly: 260		ı Ala	a Ser	- Ala	a Asp 265		e Glu	Thr	· Tyr	- Leu 270		G G L u
10	Lya	s Se	r Ar 27	g Va 5	l Thr	- Ph€	e Gir	Leu 280		: Ala	Glu	Arg	Ser 285		His	i lle
15	Pho	∋ Ty 29		n Ile	e Leu	Ser	Asn 295		: Lys	Pro	Glu	Leu 300		Glu	Met	Leu
20	Le:		e Th	r Thr	- Asn	Pro 310		Asp	Phe	Ala	Phe 315	Val	Ser	GIn	Gly	Glu 320
	ile	Th	r Va	l Pro	Ser 325		Asp	Asp	GIn	GI u 330	Glu	Leu	Met	Ala	Thr 335	Asp
25	Ser	· Ala	a Va	1 Asp 340		Leu	Gly	Phe	Thr 345		Asp	Glu	Lys	Va I 350	Ala	lle
30	Tyr	Lys	358	ı Thr	Gly	Ala	Val	Met 360	His	Tyr	Gly	Asn	Met 365	Lys	Phe	Lys
35	Gin	Lys 370		n Arg	Glu	Glu	GIn 375	Ala	Glu	Pro	Asp	Gly 380	Thr	Glu	Val	Ala
40	Asp 385	Lys	: Ala	Ala	Tyr	Leu 390	Thr	Ser	Leu	Asn	Ser 395	Ala	Asp	Leu	Leu	Lys 400
	Ser	Leu	Cys	Tyr	Pro 405	Arg	Val	Lys	Val	Gly 410	Asn	Glu	Phe	Val	Thr 415	Lys
45	Gľy	Gin	Thr	Va I 420	GIn	Gln	Val		Asn 425	Ala	Val	Gly	Ala	Leu 430	Ala	Lys

	Ala		435		ı Lys	Met	: Phe	440		Met	: Val	l Thr	Arg 445		Asn	Gln
5	Gin	Let 450		Thr	· Lys	Gln	Pro 455		: GIn	Tyr	Phe	460		Val	Leu	Asp
10 ·	l l e 465	Ala	ı Gly	Phe	Glu	11e 470		qaA q	Phe	Asn	Ser 475		Glu	GIn	Leu	Cys 480
15	lle	Asr	Phe	Thr	Asn 485		Lys	Leu	Gin	GIn 490		Phe	Asn	His	His 495	
20	Phe	Val	Leu	Glu 500	Gln	Glu	Glu	Tyr	Lys 505		Glu	Gly	lle	Glu 510	Trp	Glu
	Phe	He	Asp 515		Gly	Met	Asp	Leu 520	Ala	Ala	Cys	lle	Glu 525	Leu	He	Glu
<b>25</b>	Lys	Pro 530		Gly	He	Phe	Ser 535	ile	Leu	Glu	Glu	Glu 540	Cys	Met	Phe	Pro
30	Lys 545	Ala	Thr	Asp	Thr	Ser 550	Phe	Lys	Asn	Lys	Leu 555		Glu	Gln	His	Leu 560
35	Gly	Lys	Ser	Asn	<b>Asn</b> 565	Phe	GIn	Lys	Pro	Lys 570	Pro	Ala	Lys	Gly	Lys 575	Pro
10	Glu	Ala	His	Phe 580	Ser	Leu	Val	His	Tyr 585	Ala	Gly	Thr	Val	Asp 590	Tyr	Asn
	He	Ala	Gly 595	Trp	Leu	Asp		Asn 600	Lys	Asp	Pro	Leu	Asn 605	Glu	Thr	 Val
15	Val	Gly 610	Leu	Tyr	Gin	Lys	Ser 615	Ala	Met	Lys	Thr	Leu 620	Ala	Phe	Leu	Phe

	Se:	c Gly	y Ala	Gir	1 Thr	630		ı Ala	( Glu	Gly	635		Gly	' Lys	Lys	640
5	Gly	/ Lys	s Lys	Lys	645		Ser	Phe	Gin	Thr 650		Ser	Ala	Leu	Phe 655	
10	Glu	ı Asr	ı Leu	Asn 660	Lys	Leu	Met	Thr	Asn 665		Arg	Ser	Thr	His 670		His
15	Phe	Val	Arg 675		lle	lle	Pro	Asn 680	Glu	Thr	Lys	Thr	Pro 685		Ala	Met
20	Glu	His 690		Leu	Val	Leu	His 695		Leu	Arg	Cys	Asn 700		Val	Leu	Glu
	Gly 705	He	Arg	l le	Cys	Arg 710	Lys	Gly	Phe	Pro	Ser 715	Arg	He	Leu	Tyr	Ala 720
25	Asp	Phe	Lys	Gln	Arg 725	Tyr	Lys	Val	Leu	<b>Asn</b> 730	Ala	Ser	Ala	He	Pro 735	
30	Gly	Gln	Phe	11e 740	Asp	Ser	Lys	Lys	Ala 745	Ser	Glu	Lys	Leu	Leu 750	Gly	Ser
35	lie	Glu	lle 755	Asp	His	Thr	Gin	Tyr 760	Lys	Phe	Gly	His	Thr 765	Lys	Val	Phe
40	Phe	Lys 770	Ala	Gly	Leu	Leu	Gly 775	Thr	Leu	Glu	Glu	Met 780	Arg	Asp	Glu	Lys
	Leu 785	Ala	Gin	Leu	lle	Thr 790	Arg	Thr	GIn	Ala	lle 795	Cys	Arg	Gly	Phe	Leu 800
45	Met	Arg	Val		Phe 805	Arg	Lys	Met		Glu 810	Arg	Arg	Glu	Ser	lle 815	Phe

	Cys	: []e	e Gin	820		ı.Ile	Arg	Ala	Phe 825		: Asn	Val	Lys	His 830		Pro
5	Trp	) Met	t Lys 835		ı Tyr	Phe	Lys	840		Pro	Leu	Leu	Lys 845	Ser	Ala	Glu
10	Thr	61u 850	ı Lys )	Glu	Met	Ala	Asn 855		Lys	Glu	Glu	Phe 860		Lys	Thr	Lys
15	G1u 865		ı Leu	Ala	Lys	Thr 870		Ala	Lys	Arg	Lys 875	Glu	Leu	Glu	Glu	Lys 880
20	Met	Val	Thr	Leu	Met 885	Gin	Glu	Lys	Asn	Asp 890	Leu	<b>GI</b> n	Leu	Gin	Va I 895	Gin
	Ala	Glu	Ala	Asp 900	Ala	Leu	Ala	Asp	Ala 905	Glu	Giu	Arg	Cys	Asp 910	Gln	Leu
25	lle	Lys	Thr 915	Lys	lie	Gin	Leu	Glu 920	Ala	Lys	Ļle	Lys	Glu 925	Val	Thr	Glu
30	Arg	Ala 930	Glu	Asp	Glu	Glu	Glu 935	He	Asn	Ala	Glu	Leu 940	Thr	Ala	Lys	Lys
35	Arg 945	Lys	Leu	Glu	Asp	Giu 950	Cys	Ser	Glu	Leu	Lys 955	Lys	Asp	lle	Asp	Asp 960
40	Leu	GIú	Leu	Thr	Leu 965	Ala	Lys	Val		Lys 970	Glu	Lys	His	Ala	Thr 975	Glu
	Asn	Lys	Val	Lys 980	Asn	Leu	Thr	Glu	Glu 985	Met	Ala	Gly		Asp 990	Glu	Thr
45	lle	Ala	Lys 995	Leu	Thr	Lys	Glu	Lys 1000		Ala	Leu	Gln	Glu 100		a Hi	s Gin

	GIn	1010		Asp	Asp	Leu	GIn 1015		Glu	Glu	Asp	Lys 1020		Asn	Thr
5	Leu	Thr 1025		Ala	Lys	Thr	Lys 1030		Glu	GIn	Gln	Val 1035	Asp	Asp	Leu
10	Glu	Gly 1040		Leu	Glu	Gin	Glu 1045		Lys	Leu	Cys	Met 1050		Leu	Glu
15	Arg	Ala 1055		Arg	Lys	Leu	Glu 1060	Gly	Asp	Leu	Lys	Leu 1065	Ala	Gin	Glu
20	Ser	Thr 1070		Asp	Thr	Glu	Asn 1075		Lys	Gin	Gln	Leu 1080	Asn	Glu	Lys
	Leu	Lys 1085		Lys	Glu	Phe	Glu 1090		Ser	Asn	Leu	GIn 1095	Gly	Lys	He
25	Glu	Asp 1100		Gln	Ala	Leu	Ala 1105	Met <sub>.</sub>	Gin	Leu	Gin	Lys 1110	Lys	He	Lys
30	Glu	Leu 1115		Ala	Arg	He	Glu 1120	Glu	Leu	Glu	Glu	Glu 1125	lle	Glu	Ala
35	Glu	Arg 1130		Ser	Arg	Ala	Lys 1135	Ala	Glu	Lys	Gln	Arg 1140	Ser	Asp	Leu
40	Ser	Arg 1145	Glu	Leu	Glu	Glu	l le 1150	Ser	Glu	Arg	Leu	Glu 1155	Glu	Ala	Gly
	Gly	Ala 1160	Thr	Ser	Ala	GIn	lie 1165	Glu	Leu	Asn	Lys	Lys 1170	Arg	Glu	Ala
15	Glu	Phe 1175	<b>G</b> In	Lys	Met	Arg	Arg 1180	Asp	Leu	Glu	Giu	Ser 1185	Thr	Leu	Gin

	HIS	1190		a Thr	- Ala	Ala	1 Ala 1195		Arg	Lys	Lys	His 1200		Asp	Ser
5	Val	1205		. Leu	ı Giy	' Lys	GIn 1210		Asp	Ser	Leu	GIn 1215		Val	Lys
10	Gir	Lys 1220		ı Glu	l Lys	Glu	Lys 1225	Ser	Glu	Leu	Lys	Met 1230		lie	Asn
15	Asp	Leu 1235		Ser	Asn	Met	Glu 1240		Vai	Ser	Lys	Ala 1245		Ala	Asn
20	Phe	Glu 1250		Met	Cys	Arg	Thr 1255		Glu	Asp	Gin	Leu 1260		Glu	lle
	Lys	Thr 1265		Glu	Glu	Glu	Gin 1270		Arg	Leu	lle	Asn 1275	Glu	Leu	Ser
25	Ala	GIn 1280		Ala	Arg	Leu	His 1285		Glu	Ser	Gly	Glu 1290	Phe	Ser	Arg
30	GIn	Leu 1295		Glu	Lys	Asp	Ala 1300		Val	Ser	Gln	Leu 1305	Ser	Arg	Gly
35	Lys	GIn 1310		Phe	Thr	Gin	GIn 1315		Glu	Glu	Leu	Lys 1320	Arg	Gin	Leu
	Glu	Glu 1325	Glu	Thr	Lys	Ala	Lys 1330	Ser	Thr	Leu	Ala	His 1335	Ala	Leu	GIn
	Ser	Ala 1340	Arg	His	Asp	Cys	Asp 1345	Leu	Leu	Arg	Glu	GIn 1350	Tyr	Glu	Glu
45	Glu	Gin 1355	Glu	Ala	Lys	Ala	Glu 1360	Leu	Gln	Arg	Gly	Met 1365	Ser	Lys	Ala

	Asn	Ser 1370		Val	Ala	Gin	Trp 1375	Arg	Thr	Lys	Tyr	Glu 1380	Thr	Asp	Ala
5	He	GIn 1385	Arg	Thr	Glu	Glu	Leu 1390	<b>Gl</b> u	Glu	Ala	Lys	Lys 1395	Lys	Leu	Ala
10	GIn	Arg 1400		Gln	Asp	Ala	Glu 1405	@lu	His	Val	Glu	Ala 1410	Val	Asn	Ser
15	Lys	Cys 1415	Ala	Ser	Leu	Glu	Lys 1420	Thr	Lys	<b>G</b> In	Arg	Leu 1425	GIn	Asn	Glu
20	Val	Glu 1430	Asp	Leu	Met	He	Asp 1435	Val	Glu	Arg	Ser	Asn 1440	Ala	Ala	Cys
	lle	Ala 1445	Leu	Asp	Lys	Lys	GIn 1450	Arg	Asn	Phe	Asp	Lys 1455	Val	Leu	Ala
25	Glu	Trp 1460	Lys	Gln	Lys	Tyr	GIu 1465	Glu	Thr	GIn	Ala	Glu 1470	Leu	Glu	Ala
30	Ser	GIn 1475	Lys	Glu	Ser	Arg	Ser 1480	Leu	Ser	Thr	Glu	Leu 1485	Phe	Lys	Val
35	Lys	Asn 1490	Ala	Tyr	Glu	Glu	Ser 1495	Leu	Asp	His	Leu	Glu 1500	Thr	Leu	Lys
10		Glu 1505	Asn	Lys	Asn	Leu	GIn 1510	GIn	Glu	He	Ser	Asp 1515	Leu	Thr	Glu
	Gin	l le 1520	Ala	Glu	Gly	Gly	Lys 1525	His	lle	His	Glu	Leu 1530	Glu	Lys	Val
15	Lys	Lys 1535	Gin	Leu	Asp	His	Glu 1540	Lys	Ser	Glu	Leu	GIn 1545	Thr .	Ser	Leu

	Glu	GIU 1550		Glu	Ala	Ser	Leu 1555		His	Glu	Glu	Gly 1560		lle	Leu
5	Arg	lle 1565		Leu	Glu	Leu	Asn 1570		Val	Lys	Ser	01u 1575		Asp	Arg
10	Lys	IIe 1580		Glu	Lys	Asp	Glu 1585		Leu	Asp	Gln	Leu 1590		Arg	Asn
15	His	Leu 1595		Vaİ	Val	Glu	Ser 1600	Met	GIn	Ser	Thr	Leu 1605	Asp	Ala	Glú
20	lle	Arg 1610		Arg	Asn	Ąsp	Ala 1615	Leu	Arg	lle	Lys	Lys 1620	Lys	Met	Glu
	Gly	Asp 1625	Leu	Asn	Glu	Met	Glu 1630	lle	Gin	Leu	Asn	His 1635	Ala	Asn	Arg
25	Gln	Ala 1640	Ala	Glu	Ala	Leu	Arg 1645	Asn	Leu	Arg	Asn	Thr 1650	Gln	Gly	lle
30	Leu	Lys 1655	Asp	Thr	Gln	Leu	His 1660	Leu	Asp	Asp	Ala	l le 1665	Arg	Gly	Gln
35	Asp	Asp 1670	Leu	Lys	Glu	GIn	Leu 1675	Ala	Met	Val	Glu	Arg 1680	Arg	Ala	Asn
40	Leu	Met 1685	Gln	Ala	Glu	Val	Glu 1690	Glu	Leu	Arg	Ala	Ser 1695	Leu	Glu	Arg
	Thr	Glu 1700	Arg	Gly	Arg	Lys	Met 1705	Ala	Glu	GIn	Glu	Leu 1710	Leu	Asp	Ala
45	Ser	Glu 1715	Arg	Val	Gin	Leu	Leu 1720	His	Thr	GIn	Asn	Thr 1725	Ser	Leu	He

	ISA	1730	Ly: )	6 Ly	s Ly:	s Lei	ı Glu 173!		r Ası	o Ile	e Ser	1740		Gli	n Gly
5	Glı	ı Met 1745	GI:	u Ast	)   [ (	e Val	1 Gin 1750		u Ala	a Arg	, Asr	1758		ı Glı	ı Lya
10	Ala	1760	Lys	S Ala	ı He	• Thr	- Asp 1765		a Ala	Met	Met	: Ala 1770		Glu	ı Leu
15	Lys	Lys 1775	Gl.	ı Gin	Asp	Thr	Ser 1780		a His	Leu	Glu	Arg 1785		Lys	; Lys
20	Asn	Met 1790	Glu	ı Gin	Thr	· Val	Lys 1795		Leu	Gln	Leu	Arg 1800		Gly	Glu
	Ala	GI u 1805	Gin	Leu	Ala	Leu	Lys 1810		Gly	Lys	Lys	GIn 1815		Gln	Lys
25	Leu	Glu 1820	Ala	Arg	Vai	Arg	Glu 1825		Glu	Ser	Glu	Val 1830	Glu	Ser	Glu
30	GIn	Lys 1835	His	Asn	Val	Glu	Ala 1840		Lys	GÌy	Leu	Arg 1845	Lys	His	Glu
35	Arg	Arg 1850	Val	Lys	Glu	Leu	Thr 1855		GIn	Thr	Glu	Glu 1860	Asp	Arg	Lys
40	Asn	lle 1865	Leu	Arg	Leu	Gin	Asp 1870	Leu	Val	Asp	Lys	Leu 1875	GIn	Thr	Lys
	Val	Lys 1880	Ala	Tyr	Lys	Arg	GIn 1885	Ala	Glu	Glu	Ala	GIu 1890	Glu	Gln	Ser
<b>45</b>	Asn	Va I 1895	Asn	Leu	Ala		Phe 1900	Arg	Lys	Leu		His 1905	Glu	Leu	Glu

Glu Ala Glu Glu Arg Ala Asp lie Ala Glu Ser Gln Val Asn Lys 1910 1915 1920

Leu Arg Val Lys Ser Arg Glu Val His Thr Lys Val I le Ser Glu 1925 1930 1935

10 Glu

<210> 5
15 <211> 5925
 <212> DNA
 <213> Homo sapiens

**<400>** 5 20 atgagttctg actctgagat ggccattttt ggggaggctg ctcctttcct ccgaaagtct 60 gaaagggagc gaattgaagc ccagaacaag ccttttgatg ccaagacatc agtctttgtg 120 gtggacccta aggagtcctt tgtgaaagca acagtgcaga gcagggaagg ggggaaggtg 180 25 acagctaaga cogaagctgg agctactgta acagtgaaag atgaccaagt cttccccatg 240 aaccctccca aatatgacaa gatcgaggac atggccatga tgactcatct acacgagcct 300 30 gctgtgctgt acaacctcaa agagcgctac gcagcctgga tgatctacac ctactcaggc 360 ttgttctgtg tcactgtcaa cccctacaag tggttgccag tgtataatgc agaagtggtg 420 acagoctaco gaggoaaaaa gogocaggaa gococacoco acatottoto catototgac 480 35 aatgoctato agttoatgot gactgatogg gagaatcagt ctatottgat cacoggagaa 540 totggcgcag ggaagactgt gaacaccaag cgtgtcatcc agtactttgc aacaattgca 600 40 gttactgggg agaagaagaa ggaagaagtt acttctggca aaatgcaggg gactctggaa 660 gatcaaatca tcagtgccaa cccctactg gaggcctttg gcaacgccaa gaccgtgagg 720 aatgacaact cctctcgctt tggtaaattc atcaggatcc acttcggtac cacagggaaa 780 45 ctggcttctg ctgatattga aacatatctt ctggagaagt ctagagttac tttccagcta 840 aaggotgaaa gaagotatoa tatttttat cagatoatgt otaacaagaa gocagatota 900

	attgaaatgo	c tootgatcad	Caccaaccc	a tacgattate	g cottogtoa	g toaaggggag	960
5	atcacagtgo	c ccagcattga	a tgaccaagaa	a gagttgatgg	g ctacagatas	g tgccattgaa	1020
	attotgggot	t ttacttcage	a tgaaagagtg	g tocatotate	a agotoacage	ggctgtgatg	1080
	cattatggga	a acatgaaati	: caagcaaaa	g cagogtgagg	g agcaagctge	gccagatggc	1140
10	actgaagttg	g ctgacaaggo	agoctatoto	caaaatctga	actotgoaga	tctgctcaaa	1200
	gocototgot	: accctagggt	: caaggtcggc	aatgagtatg	tcaccaaagg	tcaaactgtg	1260
15	cagcaggtgt	: acaatgcagt	gggtgctctg	gccaaagctg	tctacgataa	gatgttcttg	1320
	tggatggtca	cccgcatcaa	ccagcagctg	gacaccaago	agcccaggca	gtacttcatt	1380
	ggggtcttgg	acattgctgg	ctttgagato	tttgatttca	acagcctgga	gcagctgtgc	1440
20	atcaacttca	ccaatgagaa	actgcaacag	: tttttcaacc	accacatgtt	cgtgctggag	1500
	caggaggagt	acaagaagga	aggcattgag	tggacgttca	ttgactttgg	gatggacctg	1560
25	gctgcctgca	tcgagctcat	cgagaagcct	atgggcatct	tctccatcct	ggaagaggag	1620
	tgcatgttcc	ccaaggcgac	agacacctcc	ttcaagaaca	agctgtatga	acaacatctt	1680
	ggaaaatcca	ataacttcca	gaagcccaag	cctgccaaag	gcaagcctga	ggcccacttc	1740
30	tctttgattc	actatgctgg	caccgtggac	tacaacattg	ccggctggct	tgacaagaac	1800
	aaggaccccc	tgaatgagac	tgtggtgggg	ctgtaccaga	agtctgcaat	gaagactctg	1860
35	gctctcctct	ttgttggggc	aacgggagcg	gaagcagagg	ctggcggtgg	aaagaaaggt	1920
	ggtaagaaga	agggttcttc	tttccagact	gtgtcggctc	tcttcaggga	gaatttgaat	1980
	aagctgatga	ccaacttgag	gagcactcac	cccactttg	tgcggtgcat	catccccaat	2040
10	gaaactaaaa	ctcctggtgc	catggagcat	gagcttgtcc	tgcatcagct	gaggtgtaac	2100
	ggtgtgctgg	aaggcatccg	catctgcagg	aaaggcttcc	caagcagaat	cctttatgca	2160
5	gacttcaaac	agagatacaa	ggtgttaaat	gcaagtgcta	tccctgaagg	acaattcatc	2220
	gatagcaaga	aggcttcaga	gaagctcctg	gggtccattg	acattgacca	cacccagtat	2280
	aaatttggtc	acaccaaggt	ctttttcaaa	gctggtcttc	terrectect	agaggagatg	2340

	cgagatgaga	agctggccca	gctgattacc	cgaacccagg	ccatgtgcag	agggttcttg	2400
5	gcaagagtgg	agtaccagaa	aatggtggaa	agaagagagt	ccatcttotg	catccagtac	2460
	aatgtccgtg	ccttcatgaa	tgtcaagcac	tggccctgga	tgaagctgta	tttcaagatc	2520
	aaacccctcc	tcaaaagtgc	agagacagag	aaggagatgg	ccaacatgaa	ggaagaattt	2580
10	gagaaaacca	aagaagagct	ggctaagacc	gaggcaaaaa	ggaaagagct	ggaagaaaaa	2640
	atggtgactc	tgatgcaaga	aaaaaatgac	ttgcaactcc	aggttcaagc	tgaagctgac	2700
15	agcttggctg	atgcagagga	aaggtgtgac	cagctaatca	aaaccaaaat	ccagctagaa	2760
	gccaaaatca	aagaggtgac	tgagagagct	gaggatgagg	aagagatcaa	tgctgagctg	2820
	acagccaaga	agaggaaact	ggaggatgaa	tgttcagaac	tcaagaaaga	cattgatgac	2880
20	cttgagctga	cactggccaa	ggttgagaag	gagaaacatg	ccacagaaaa	caaggtgaaa	2940
	aacctcacag	aagagatggc	gggtctggat	gaaaccattg	ctaagctgac	caaggagaag	3000
25	aaggototoo	aggaggccca	ccagcagacc	ctggatgacc	tgcaggcaga	ggaggacaaa	3060
	gtcaacaccc	tgaccaaagc	taaaatcaaa	cttgaacaac	aagtggatga	tcttgaagga	3120
	tctttggaac	aagaaaagaa	aatccggatg	gatctagaaa	gagcaaagag	aaaactagag	3180
30	ggagacctaa	aattggctca	agaatccgca	atggatatag	aaaatgacaa	acaacaactt	3240
	gatgaaaagc	ttaaaaagaa	agagtttgaa	atgagcggtc	tgcaaagcaa	gattgaagat	3300
35	gaacaagccc	ttggtatgca	gctgcagaag	aaaatcaagg	agttacaagc	ccgcattgag	3360
-	gagctggagg	aggaaatcga	ggcagagcgg	gcctcccggg	ccaaagcaga	gaagcagcgc	3420
	totgatotot	cccgggagct	ggaggagatc	agtgagaggc	tggaagaagc	cggtggggcc	3480
40	acctcggccc	agattgagat	gaacaagaag	cgggaagctg	agttccagaa	aatgogoagg	3540
	gacctggagg	aggccaccct	acagcatgag	gccacggcgg	ccaccctgag	gaagaagcat	3600
45	gcagatagtg	tggccgagct	tggggagcag	attgacaacc	tgcagcgagt	gaagcagaag	3660
-~	ctggagaagg	agaagagtga	gatgaagatg	gagatcgatg	accttgctag	taacatggag	3720
	actgtctcca	aagccaaggg	aaaccttgaa	aagatgtgcc	gcgctctaga	agatcaactg	3780

	agtgaaatta	a agaccaagg	a agaggagca	g cagoggotga	tcaatgacci	: cacagcacag	3840
5	agagogogo	tgcaaacag	a atcaggtga	a tattcacgco	agotagatga	aaaggacaca	3900
J	ctagtttcac	agctctcga	g gggcaaacas	gootttacto	aacagattga	ggaactgaaa	3960
•	aggcaacttg	s aagaggaga	t aaaggocaag	g agtgocotgg	cacatgocot	gcagtcctcc	4020
10	cgccatgact	gtgacctgc	t gcgggaacag	g tatgaggagg	agcaggaagc	caaggccgag	4080
	ctacagagag	caatgtccaa	a ggccaacagt	gaggttgcco	agtggaggac	caaatatgag	4140
15	acagatgcca	tocagogoad	agaggagctg	gaggaggoca	agaagaagct	ggctcagcgt	4200
	ctgcaggatg	ctgaggaaca	tgtagaagct	gtgaatgcca	aatgtgcttc	ccttgagaag	4260
	acgaagcaga	ggctccagaa	ı tgaagttgag	gacctcatga	ttgatgttga	gaggacaaat	4320
20	gctgcctgtg	ccgccctgga	caaaaagcaa	aggaactttg	ataagatoot	ggcagaatgg	4380
	aaacagaagt	gtgaagaaac	tcatgotgaa	cttgaagctt	ctcaaaagga	atcccgctca	4440
25	ctcagcacag	aactatttaa	gattaagaat	gcttatgagg	aatctttaga	ccaacttgaa	4500
	accttgaaac	gggaaaataa	gaatctgcag	caggagattt	ctgatctcac	tgaacagatt	4560
	gcagaaggag	gaaagcgcat	ccatgaactg	gaaaaaataa	agaagcaagt	tgagcaagaa	4620
30	aagtotgaac	ttcaggctgc	cttagaggag	gcagaggcat	ctcttgaaca	tgaagagga	4680
	aagatcctgc	gcatccagct	tgagttgaac	caagtcaagt	ctgaggttga	taggaaaatt	4740
35 .	gotgaaaaag	atgaggaaat	tgaccagatg	aagagaaacc	acattagaat	cgtggagtcc	4800
	atgcagagca	cactggatgc	tgagatcagg	agcaggaatg	atgccattag	gctcaagaag	4860
	aagatggagg	gagacctcaa	tgaaatggaa	atccagctga	accatgccaa	ccgcatggct	4920
10	gctgaggccc	tgaggaacta	taggaacacc	caagccatcc	tcaaggatac	ccagctccac	4980 °
	ctagatgatg	ctctccggag	ccaagaggac	ctgaaggaac	agctggctat	ggtggagcgc	5040
15	agagccaacc	tgctgcaggc	tgagatcgag	gaactacgag	ccactctgga	acagacggag	5100
	aggagcagga	aaatcgcaga	acaggagete	ctggatgcca	gtgaacgtgt	toagctcctg	5160
	cacacccaga	acaccagcct	gatcaacacc	aagaagaagc	tggagacaga	catttcccaa	5220

	atccagggag agatggaaga catcatccag gaagcccgca atgcagaaga gaaggccaag	5280
5	aaggocatca ctgatgotgo catgatggot gaggagotga agaaggaaca ggacaccago	5340
J	goccatotgg agoggatgaa gaagaacttg gaacagacgg tgaaggacct goagcatogt	5400
	ctggatgagg ctgagcagct ggccctgaag ggtgggaaga agcagatcca gaaactggag	5460
10	gccagggttc gtgaacttga aggtgaagtt gaaagtgaac agaagcgcaa tgttgaagct	5520
	gtcaagggtc tacgcaaaca tgagagaaaa gtgaaggaac tcacttacca aactgaggaa	5580
15	gaccgcaaga atattotoag gotgcaggac otggtggaca agotgcaago aaaggtgaaa	5640
	toctacaaga gacaagotga agaagoggag gaacaatoca acgtcaacot otocaaatto	5700
	cggaggatcc agcacgagct ggaggaggcc gaggaaaggg ctgacattgc tgagtcccag	5760
20	gtcaacaagc tgagggtgaa gagcagggag gttcacacaa aaatcataag tgaagagtaa	5820
	tttatctaac tgctgaaagg tgaccaaaga aatgcacaaa atgtgaaaat ctttgtcact	5880
25	ccattttgta cttatgactt ttggagataa aaaatttatc tgcca	5925
30	<210> 6 <211> 1939 <212> PRT <213> Homo sapiens	·
	<400> 6	
35	Met Ser Ser Asp Ser Glu Met Ala IIe Phe Gly Glu Ala Ala Pro Phe 1 5 10 15	
40	Leu Arg Lys Ser Glu Arg Glu Arg IIe Glu Ala Gln Asn Lys Pro Phe 20 25 30	
	Asp Ala Lys Thr Ser Vai Phe Vai Vai Asp Pro Lys Glu Ser Phe Vai 35 40 45	
45		

Lys Ala Thr Val Gin Ser Arg Glu Gly Gly Lys Val Thr Ala Lys Thr

60

55

	61 t	i Ala	a Gly	/ Ala	a Thr	70	l Thr	· Val	Lys	s Asp	Asp 75	GIr	\Val	Phe	Pro	Met 80
5	Asr	n Pro	o Pro	) Lys	Tyr 85	· Asp	Lys	lle	Glu	Asp 90	Met	Ala	Met	: Met	: Thr : 95	His
10	Leu	ı His	s Glu	Pro 100		val	Leu	ı Tyr	Asn 105	Leu	Lys	Glu	Arg	Tyr 110		Ala
15	Trp	) Met	: lle 115		Thr	Tyr	· Ser	Gly 120		Phe	Cys	Val	Thr 125		Asn	Pro
20	Tyr	Lys 130		Leu	Pro	Val	Tyr 135		Ala	Glu	Val	Va I 140		Ala	Tyr	Arg
	Giy 145		Lys	Arg	Gln	Glu 150		Pro	Pro	His	l le 155	Phe	Ser	ile		Asp 160
25	Asn	Ala	Tyr	GIn	Phe 165	Met	Leu	Thr	Asp	Arg 170	Glu	Asn	Gln	Ser	l i e 175	Leu
30	lle	Thr	Gly	Glu 180	Ser	Gly	Ala	Gly	Lys 185	Thr	Vai	Asn	Thr	Lys 190	Arg	Val
35	lle	Gln	Tyr 195	Phe	Ala	Thr	lie	Ala 200	Val	Thr	Gly	Glu	Lys 205	Lys	Lys	Glu
10	Glu	Va l 210	Thr	Ser	Gly	Lys	Met 215	GIn	Gly	Thr	Leu	Glu 220	Asp	Gln	lle	He
	Ser 225	Ala	Asn	Pro	Leu	Leu 230	Glu	Ala	Phe	Gly	Asn 235	Ala	Lys	Thr	Val	Arg 240
5	Asn	Asp	Asn	Ser	Ser 245	Arg	Phe	Gly	Lys	Phe 250	He	Arg	He	His	Phe	Gly

	ınr	' Ihr	Gly	260		Ala	Ser	Ala	Asp 265		Glu	Thr	Tyr	Leu 270		Glu
5	Lys	Ser	• <b>A</b> rg 275		Thr	Phe	Gin	Leu 280	Lys	Ala	Glu	Arg	Ser 285		His	He
10	Phe	Tyr 290		lle	Met	Ser	Asn 295		Lys	Pro	Asp	Leu 300		Glu	Met	Leu
15	Leu 305		Thr	Thr	Asn	Pro 310	Tyr	Asp	Tyr	Ala	Phe 315	Val	Ser	Gln	Gly	Glu 320
20	lle	Thr	Val	Pro	Ser 325	lle	Asp	Asp	Gin	Giu 330	Glu	Leu	Met	Ala	Thr 335	Asp
	Ser	Ala	lle	G1u 340	He	Leu	Gly	Phe	Thr 345	Ser	Asp	Glu	Arg	Va I 350	Ser	lle
25	Tyr	Lys	Leu 355	Thr	Gly	Ala	Val	Met 360	His	Tyr	Gly	Asn	Met 365	Lys	Phe	Lys
30	GIn	Lys 370	GIn	Arg	Glu	Glu	GIn 375	Ala	Glu	Pro	Asp	Gly 380	Thr	Glu	Val	Ala
35	Asp 385	Lys	Ala	Ala	Tyr	Leu 390	GIn	Asn	Leu-	Asn	Ser 395	Ala	Asp	Leu	Leu	Lys 400
10	Ala	Leu	Cys	Tyr	Pro 405	Arg	Val	Lys	Val	Gly 410	Asn	Glu	Tyr	Vai	Thr 415	Lys
	Gly	Gln	Thr	Va I 420	Gln	<b>G</b> In	Val	Tyr	Asn 425	Ala	Val	Gly	Ala	Leu 430	Ala	Lys
15	Ala	Val	Tyr 435	Asp	Lys	Met	Phe	Leu 440	Trp	Met	Val	Thr	Arg 445	He	Asn	GIn

	Gir	1 Let 450		Thr	· Lys	Gin	Pro 455		Gln	Tyr	· Phe	11e 460		Val	Leu	Asp
	l l e 465		Gly	Phe	Glu	11e 470		Asp	Phe	Asn	Ser 475		Glu	Gin	Leu	Cys 480
10	He	Asn	Phe	Thr	Asn 485		Lys	Leu	Ğln	G1n 490		Phe	Asn	His	His 495	
15	Phe	Val	Leu	Glu 500	GIn	Glu	Glu	Tyr	Lys 505		Glu	Gly	He	Glu 510		Thr
20	Phe	He	Asp 515	Phe	Gly	Met	Asp	Leu 520		Ala	Cys	ile	Glu 525		lle	Glu
	Lys	Pro 530		Gly	lle	Phe	Ser 535	ile	Leu	Glu	Glu	Glu 540	Cys	Met	Phe	Pro
25	Lys 545	Ala	Thr	Asp	Thr	Ser 550	Phe	Lys	Asn	Lys	Leu 555	Tyr	Glu	Gin	His	Leu 560
30	Gly	Lys	Ser ·	Asn	Asn 565	Phe	Gin	Lys	Pro	Lys 570	Pro	Ala	Lys	Gly	Lys 575	Pro
35	Glu	Ala	His	Phe 580	Ser	Leu	He	His	Tyr 585	Ala	Gly	Thr	Val	Asp 590	Tyr	Asn
40	He	Ala	Gly 595	Trp	Leu	Asp	Lys	Asn 600	Lys	Asp	Pro	Leu	Asn 605	Glu	Thr	Val
	Val	Gly 610	Leu	Tyr	Gln		Ser 615	Ala	Met	Lys	Thr	Leu 620	Ala	Leu	Leu	Phe
45	Va I 625	Gly	Ala	Thr	Gly	Ala 630	Glu	Ala	Glu	Ala	Gly 635	Gly	Gly	Lys	Lys	Gly 640

	GI	y Ly	's Ly:	s Ly:	64!	y Sei 5	r Sei	· Pho	e Glr	1 Thr 650		Sei	^ Ala	i Lei	658	
5	Glu	aA u	n Lei	18A U 186	n Lys )	s Lei	J Met	: Thi	- Asr 665		ı Arg	; Ser	- Thr	His 670		) His
10	Phe	e Va	l Arg 675	g Cya	s He	) He	Pro	Asr 680		ı Thr	· Lys	Thr	Pro 685		Ala	ı Met
15	Glu	Hi:	s Glu O	ı Let	ı Val	Leu	ı His 695		Leu	ı Arg	Cys	Asr 700		Val	Leu	ı Glu
20	Giy 705	/ [](	e Arg	; lle	Cys	710		Gly	Phe	Pro	Ser 715		lle	Leu	Tyr	Ala 720
	Asp	Phe	) Lys	Gin	Arg 725		Lys	Val	Leu	Asn 730		Ser	Ala	He	Pro 735	
25	Gly	Gir	n Phe	11e 740		Ser	Lys	Lys	Ala 745		Glu	Lys	Leu	Leu 750	Gly	Ser
30	lle	Asp	755	Asp	His	Thr	Gin	Tyr 760	Lys	Phe	Gly	His	Thr 765	Lys	Val	Phe
35	Phe	Lys 770	Ala	Gly	Leu	Leu	Gly 775	Leu	Leu	Glu	Glu	Met 780	Arg	Asp	Glu	Lys
40	Leu 785	Ala	Gin	Leu	ile	Thr 790	Arg	Thr	GIn	Ala	Met 795	Cys	Arg	Gly	Phe	Leu 800
	Ala	Arg	Val	Glu	Tyr 805	GIn	Lys	Met	Val	Glu 810	Arg	Arg	Glu		l le 815	Phe
45	Сув	lle	Gin	Tyr 820	Asn	Val	Arg		Phe 825	Met	Asn	Val	Lys	His 830	Trp	Pro

	Trp	Me1	t Lys 835		Tyr	Phe	Lys	840		Pro	Leu	Leu	Lys 845		Ala	Glu
5	Thr	G1:	ı Lys )	Glu	Met	Ala	Asn 855		Lys	Glu	Glu	Phe 860	Glu	Lys	Thr	Lys
10	Glu 865	Glu	ı Leu	Ala	Lys	Thr 870		Ala	Lys	Arg	Lys 875	Glu	Leu	Glu	Glu	Lys 880
15	Met	Val	Thr	Leu	Met 885	Gin	Glu	Lys	Asn	Asp 890	Leu	Gin	Leu	Gln	Va I 895	Gln
20	Ala	Glu	Ala	Asp 900	Ser	Leu	Ala	Asp	A1a 905	Glu	Giu	Arg	Cys	<b>Asp</b> 910	Gin	Leu
	lle	Lys	Thr 915	Lys	He	GIn	Leu	Glu 920	Ala	Lys	He	Lys	Glu 925	Val	Thr	Glu
25	Arg	Ala 930	Glu	Asp	Glu	Glu	Glu 935	lle	Asn	Ala	Glu	Leu 940	Thr	Ala	Lys	Lys
30	Arg 945	Lys	Leu	Glu	Asp	Glu 950	Cys	Ser	Glu	Leu	Lys 955	Lys	Asp	He	Asp	Asp 960
35	Leu	Glu	Leu		Leu 965	Ala	Lys	Val		Lys 970	Glu	Lys	His		Thr 975	Glu
40	Asn	Lys	Val	Lys 980	Asn	Leu	Thr		Glu 985	Met	Ala	Gly		Asp 990	Glu	Thr
	lle .	Ala	Lys 995	Leu	Thr	Lys	Glu	Lys 1000		Ala	Leu	Gin	Glu 100		a Hi	s Glr
45	GIn :	Thr 1010	Leu )	Asp	Asp	Leu	GIn 101		a GI	u GI	u Ası	p Ly. 10:		al A	sn T	hr

	Le	u ini 102	- Ly 25	'S AI	a Ly	s II	e Lys 103	Le <sub>i</sub> 0	u Gli	u Gir	a Gir	1035		Asp	Leu
5	GI	u Gly 104	/ Se 10	r Le	u Gi	u Gli	n Glu 104	Ly:	s Lya	s lie	Arg	Met 1050		Leu	ı Glu
10	Arį	g Ala 105	ı Ly. 5	s Ar	g Ly:	s Lei	ı Glu 1060	Gl)	/ Asp	Leu	Lys	Leu 1065		Gin	Glu
15	Ser	- Ala 107	. Me <sup>.</sup> O	t Ası	o He	e Glu	J Asn 1075	Asp	) Lys	G G I n		Leu 1080		Glu	Lys
20	Leu	l Lys 108	Lys 5	s Lys	s Glu	Phe	01u 1090		: Ser	Gly	Leu	GIn 1095		Lys	He
	Glu	110	Glu D	ı Gir	n Ala	Leu	Gly 1105		Gin	Leu	Gin	Lys 1110		lle	Lys
25	Glu	Leu 111	G1r	n Ala	Arg	lle	Glu 1120		Leu	Glu	Glu	Glu 1125	He	Glu	Ala
30	Glu	Arg 113(	Ala	Ser	Arg	Ala	Lys 1135		Glu	Lys	Gln	Arg 1140	Ser	Asp	Leu
35	Ser	Arg 1145	Glu	Leu	Glu	Glu	ile 1150	Ser	Glu	Arg	Leu	Glu 1155	Glu	Ala	Gly
40	Gly	Ala 1160	Thr	Ser	Ala	GIn	lle 1165		Met	Asn		Lys 1170	Arg	Glu	Ala
	Glu	Phe 1175	Gln	Lys	Met	Arg	Arg 1180	Asp	Leu	Glu		Ala 1185	Thr 1	Leu	GIn
45	His	Glu 1190	Ala	Thr	Ala	Ala	Thr 1195	Leu	Arg	Lys		His 1200	Ala <i>i</i>	Asp (	Ser

	Va	1 Ala 1205		i Leu	ı Giy	, Glu	1210	E	e Asp	Asn	Leu	1215		'Vai	Lys
5	Glr	n Lys 1220	Leu )	ı Glu	ı Lys	Glu	Lys 1225		- Glu	Met	Lys	Met 1230		lle	Asp
10	Asp	Leu 1235	Ala ;	Ser	Asn	Met	Glu 1240		·Val	Ser	Lys	Ala 1245		Gly	' Asn
15	Leu	ı Glu 1250	Lys	Met	Cys	Arg	Ala 1255		Glu	Asp	GIn	Leu 1260		Glu	lle
20	Lys	Thr 1265		Glu	Glu	Glu	GIn 1270		Arg	Leu	lle	Asn 1275		Leu	Thr
	Ala	Gin 1280	Arg	Ala	Arg	Leu	Gln 1285		Glu	Ser	Gly	Glu 1290		Ser	Arg
25	Gin	Leu 1295		Glu	Lys	Asp	Thr 1300		Val	Ser	Gin	Leu 1305	Ser	Arg	Gly
30	Lys	Gin 1310	Ala	Phe	Thr	Gin	GIn 1315		Glu	Glu	Leu	Lys 1320	Arg	Gin	Leu
35	Glu	Glu 1325	Glu	He	Lys	Ala	Lys 1330		Ala	Leu	Ala	His 1335	Ala	Leu	Gin
40	Ser	Ser 1340	Arg	His	Asp	Cys	Asp 1345	Leu	Leu	Arg	Glu	GIn 1350	Tyr	Glu	Glu
	Glu	GIn 1355	Glu	Ala	Lys	Ala	Glu 1360		Gln	Arg	Ala	Met 1365	Ser	Lys	Ala
45	Asn	Ser 1370	Glu	Val	Ala		Trp 1375	Arg	Thr	Lys	Tyr	Glu 1380	Thr	Asp	Ala

	110	9 GIN 138	Ari 5	g Ihi	r Gli	ı Git	139(	Gli )	ı Glu	ı Ala	ı Lys	Lys 1395		Leu	Ala
5	Gli	n Arg 140	Lei 0	ılB ı	n Asp	Ala	a Glu 1405		His	: Val	Glu	Ala 1410		Asn	Ala
10	Lys	6 Cys 141	A18	a Ser	· Leu	Glu	Lys 1420		· Lys	6In	Arg	Leu 1425		Asn	Glu
15	Val	Glu 1430	Asp )	Leu	ı Met	He	Asp 1435		Glu	Arg	Thr	Asn 1440		Ala	Cys
20	Ala	1445	Leu ;	Asp	Lys	Lys	GIn 1450		Asn	Phe	Asp	Lys 1455		Leu	Ala
	Glu	Trp 1460	Lys )	GIn	Lys	Cys	Glu 1465		Thr	His	Ala	Glu 1470	Leu	Glu	Ala
25	Ser	GIn 1475		Glu	Ser	Arg	Ser 1480		Ser	Thr	Glu	Leu 1485	Phe	Lys	lle
30	Lys	Asn 1490	Ala	Tyr	Glu	Glu	Ser 1495	Leu	Asp	Gln	Leu	Glu 1500	Thr	Leu	Lys
35	Arg	Glu 1505		Lys	Asn	Leu	GIn 1510	GIn	Glu	He	Ser	Asp 1515	Leu	Thr	Glu
40	GIn	lle 1520	Ala	Glu	Gly	Gly	Lys 1525	Arg	He	His		Leu 1530	Glu	Lys	lle
	Lys	Lys 1535	Gln	Val	Glu	Gin	Glu 1540	Lys	Ser	Glu		GIn 1545	Ala .	Alal	Leu
45	Glu	Glu 1550	Ala	Glu	Ala		Leu 1555	Glu	His	Glu (		Gly 1560	Lys	lle I	_eu

	Arg	1565 1565		1 Leu	ı Glu	i Leu	ı Asn 1570		Val	Lys	Ser	Glu 1575		Asp	Arg
5	Lys	11e 1580	Ala	a Glu	ı Lys	Asp	Glu 1585		He	Asp	GIn	Met 1590		Arg	Asn
10.	His	lle 1595		; lle	Val	Glu	Ser 1600		Gin	Ser	Thr	Leu 1605		Ala	Glu
15	lle	Arg 1610		· Arg	: Asn	Asp	Ala 1615		Arg	Leu	Lys	Lys 1620		Met	Glu
20	Ġly	Asp 1625		Asn	Glu	Met	Glu 1630		GIn	Leu	Asn	His 1635	Ala	Asn	Arg
	Met	Ala 1640		Glu	Ala	Leu	Arg 1645		Tyr	Arg	Asn	Thr 1650	Gin	Ala	lle
25	Leu	Lys 1655		Thr	Gin	Leu	His 1660	Leu	Asp	Asp	Ala	Leu 1665	Arg	Ser	Gin
30	Glu	Asp 1670		Lys	Glu	Gln	Leu 1675	Ala	Met	Val	Glu	Arg 1680	Arg	Ala	Asn
35	Leu	Leu 1685	Gin	Ala	Glu	lle	Glu 1690	Giu	Leu	Arg	Ala	Thr 1695	Leu	Glu	Gln
40	Thr	Glu 1700	Arg	Ser	Arg	Lys	lle 1705	Ala	Glu	Gin	Glu	Leu 1710	Leu	Asp	Ala
	Ser	Glu 1715	Arg	Val	GIn	Leu	Leu 1720	His	Thr	GIn	Asn	Thr 1725	Ser	Leu	lle
45	Asn	Thr 1730		Lys	Lys	Leu	Giu 1735	Thr	Asp	lle	Ser	GIn 1740	He	Gin	Gly

		Glu	Met 1745	Glu	. Asp	lle	lle	01n 1750	Glu	Ala	Arg	Asn	Ala 1755		Glu	u Lys
	5	Ala	Lys 1760	Lys	Ala	ılle	Thr	Asp 1765		Ala	Met	Met	Ala 1770		Glu	u Leu
	10	Lys	Lys 1775	Glu	Gin	Asp	Thr	Ser 1780	Ala	His	Leu	Glu	Arg 1785	Met	Lys	s Lys
	15	Asn	Leu 1790	Glu	GIn	Thr	Val	Lys 1795	Asp	Leu	GIn	His	Arg 1800	Leu	Asp	p Glu
	20	Ala	Glu 1805	GIn	Leu	Ala	Leu	Lys 1810	Gly	Gly	Lys	Lys	GIn 1815	He	Gin	n Lys
		Leu	Glu 1820		Arg	Val	Arg	Glu 1825	Leu	Glu	Gly	Glu	Val 1830	Glu	Ser	r Glu
٠.	25	Gin	Lys 1835		Asn	Val	Glu	Ala 1840	Val	Lys	Gly	Leu	Arg 1845	Lys	His	s Glu
	30		Lys 1850	Val	Lys	Glu	Leu	Thr 1855	Tyr	GIn	Thr	Glu	Glu 1860	Asp	Arg	; Lys
	35	Asn	IIe 1865	Leu	Arg	Leu		Asp 1870	Leu	Val	Asp	Lys	Leu 1875	Gln	Ala	ı Lys
	40	Val	Lys 1880	Ser	Tyr	Lys		GIn 1885	Ala.	Glu	Glu	Ala	Glu 1890	Glu	Gln	s Ser
			Va I 1895	Asn	Leu	Ser		Phe 1900	Arg .	Arg	He		His 1905	Glu	Leu	Glu
	45		Ala 1910	Glu	Glu	Arg		Asp 1915	lle i	Ala (	Glu		Gin '	Val	Asn	Lys

Leu Arg Val Lys Ser Arg Glu Val His Thr Lys 11e 11e Ser Glu

	1925 1930 1935	
5	Glu	
10	<210> 7 <211> 2633 <212> DNA <213> Homo sapiens	
15	<220> <221> CDS <222> (38) (2584) <223>	
20	<400> 7 ccgcggcaag aacatccctc ccagccagca gattaca atg ctg caa act aag gat Met Leu Gin Thr Lys Asp 1 5	55
25	ctc atc tgg act ttg ttt ttc ctg gga act gca gtt tct ctg cag gtg Leu lie Trp Thr Leu Phe Phe Leu Gly Thr Ala Val Ser Leu Gin Val 10 15 20	103
30	gat att gtt ccc agc cag ggg gag atc agc gtt gga gag tcc aaa ttc Asp lle Val Pro Ser Gin Gly Glu lle Ser Val Gly Glu Ser Lys Phe 25 30 35	151
35	ttc tta tgc caa gtg gca gga gat gcc aaa gat aaa gac atc tcc tgg Phe Leu Cys Gin Val Ala Giy Asp Ala Lys Asp Lys Asp lie Ser Trp 40 45 50	199
40	ttc tcc ccc aat gga gaa aag ctc acc cca aac cag cag cgg atc tca Phe Ser Pro Asn Gly Glu Lys Leu Thr Pro Asn Gln Gln Arg IIe Ser 55 60 65 70	247
••	gtg gtg tgg aat gat gat tcc tcc tcc acc ctc acc atc tat aac gcc Val Val Trp Asn Asp Asp Ser Ser Ser Thr Leu Thr ile Tyr Asn Ala 75 80 85	295
45	aac atc gac gac ggc att tac aag tgt gtg gtt aca ggc gag gat Asn lie Asp Asp Ala Gly lie Tyr Lys Cys Val Val Thr Gly Glu Asp 90 95 100	343

				Ser					Asn					Gin		ctc Leu	391	
5	atg Met	tto Phe	aag Lys	aat Asn	gcg Ala	cca Pro	acc Thr 125	Pro	cag Gin	gag Glu	tto Phe	cgg Arg 130	Glu	ggg Gly	gaa Glu	gat Asp	439	
10		Val	att lie				Val					Pro					487	
15			cac His													Phe	535	
20	ata Ile	gtc Val	ctg Leu	tcc Ser 170	aac Asn	aac Asn	tac Tyr	ctg Leu	cag Gin 175	atc lle	cgg Arg	ggc Gly	atc ile	aag Lys 180	aaa Lys	aca Thr	583	
	gat Asp	gag Glu	ggc Gly 185	act Thr	tat Tyr	cgc Arg	tgt Cys	gag Glu 190	ggc Gly	aga Arg	atc He	ctg Leu	gca Ala 195	cgg Arg	ggg Gly	gag Glu	631	
25	atc He	aac Asn 200	ttc Phe	aag Lys	gac Asp	att He	cag Gln 205	gtc Val	att He	gtg Val	aat Asn	gtg Val 210	cca Pro	cct Pro	acc Thr	atc ile	679	
30			agg Arg														727	
35	gtc Val	acc Thr	ctg Leu	Val	tgc Cys 235	gat Asp	gcc Ala	gaa Glu	cgg Arg	ttc Phe 240	cca Pro	gag Glu	ccc Pro	acc Thr	atg Met 245	agc Ser	775	
40	tgg Trp	aca Thr	aag Lys	gat Asp 250	ggg Gly	gaa Glu	cag Gin	lle	gag Glu 255	caa Gin	gag Glu	gaa Glu	gac Asp	gat Asp 260	gag Glu	aag Lys	823	
	tac Tyr	He	tto Phe 265	agc Ser	gac Asp	gat Asp	agt Ser	tcc Ser 270	cag Gin	ctg Leu	acc Thr	atc He	aaa Lys 275	aag Lys	gtg Val	gat Asp	871	
15	Lys	aac Asn 280	gac Asp	gag Glu	gct Ala	Glu	tac Tyr 285	atc lle	tgc Cys	att lle	Ala	gag Glu 290	aac Asn	aag Lys	gct Ala	ggc Gly	919	

		Gir	gat Asp									Ala				atc lle 310	967
5			gta Val			GIn					Leu					Thr	1015
10			tgt Cys		Ala					He					Trp		1063
15			acc Thr 345														1111
20			gtg Val														1159
2,0			tac Tyr														1207
25			cag G1n														1255
30			cag GIn					Val									1303
35	aac Asn	atc ile	acc Thr 425	tgc Cys	gag Glu	gta Val	Phe	gcc Ala 430	tat Tyr	ccc Pro	agt Ser	gcc Ala	acg Thr 435	atc He	tca Ser	tgg Trp	1351
40	Phe		gat Asp			Leu											1399
			aac Asn		Pro												1447
<b>4</b> 5	gag Glu			Phe					Cys								1495

					Glu					Gin					Ser	tca Ser	1543	•
5				Asp										Gln		cag Gin	1591	
10													Leu			aaa Lys	1639	
15						gtg Val 540						His					1687	
20						agc Ser											1735	
	aag Lys					tac Tyr											1783	
25	ggg																1831	
30		Gly 600	Glu	Pro	Ser	Ala	Pro 605	Lys	Leu	Glu	Gly	GIn 610	Met	Gly	Glu	Asp	1879	
35	gga Gly 615	aac Asn	tct Ser	att lle	aaa Lys	gtg Val 620	aac Asn	ctg Leu	atc Ile	aag Lys	cag GIn 625	gat Asp	gac Asp	ggc Gly	ggc Gly	tcc Ser 630	1 <b>927</b>	
40	Pro			His										Ser			1975	
	aaa d Lys i		Glu					Ser					Val				2023	
45	tcc c Ser L	_eu /					Glu '					Val					2071	

			Gly					Ala					Arg			gcc Ala	2119
5		Pro					Ala					Thr				agc Ser 710	2167
10						Gly										ctg Leu	2215
15					He				ttc Phe 735	Leu					Leu		2263
20				Ala					gga Gly								2311
									gcc Ala								2359
25									acg Thr								2407
30									ccc Pro							acg Thr	2455
35									gca Ala 815							aca Thr	2503
40									gtc Val								2551
									aaa Lys		tga	tggg	tgaa	iga (	gaacc	gagca	2604
45	aaga	itcaa	aa t	aaaa	agte	a ca	cago	agc									2633

	<b>&lt;2</b> 1	2>	PRT													
5		13>		sat	oiens	•										
J			ı Gin	Thr	Lys 5	Asp	Leu	ılle	Trp	. Thr 10	Leu	Phe	Phe	Leu	15	Thr
10	Ala	\Val	Ser	Leu 20	ı Gin	Val	Asp	lle	Va I 25	Pro	Ser	GIn	Gly	G I u 30	ılle	Ser
15	Val	Gly	' Glu 35	Ser	<sup>,</sup> Lys	Phe	Phe	Leu 40	Cys	Gln	Val	Ala	Gly 45	Asp	Ala	Lys
20	Asp	Lys 50	: Asp	He	Ser	Trp	Phe 55	Ser	' Pro	Asn	Gly	Glu 60	Lys	Leu	Thr	Pro
25	Asn 65	Gin	Gln	Arg	lle	Ser 70	Val	Vai	Trp	Asn	Asp 75	Asp	Ser	Ser	Ser	Thr 80
	Leu	Thr	lle	Tyr	Asn 85	Ala	Asn	lle	Asp	Asp 90	Ala	Gly	He	Tyr	Lys 95	Cys
30	Va I	Val	Thr	Gly 100	Glu	Asp	Gly	Ser	Glu 105	Ser	Glu	Ala	Thr	Val 110	Asn	Val
35	Lys	lle	Phe 115	Gin	Lys	Leu	Met	Phe 120	Lys	Asn	Ala	Pro	Thr 125	Pro	Gin	Glu
40	Phe	Arg 130	Glu	Gly	Glu	Asp	Ala 135	Val	He	Vai	Cys	Asp 140	Val	Val	Ser	Ser
45	Leu 145	Pro	Pro	Thr	lle	lle 150	Trp	Lys	His	Lys	Gly 155	Arg	Asp	Val	lle	Leu 160
	Lys	Lys	Asp	Vai	Arg 165	Phe	He	Val	Leu	Ser 170	Asn	Asn	Tyr	Leu	GIn 175	He

5	Arg	g Gly	y Ile	180		Thr	· Asp	Glu	Gly 185		Tyr	Arg	Cys	190		Arg
	He	Let	ı Ala 195		Gly	Glu	lle	Asn 200		Lys	Asp	lle	GIn 205		lle	Val
10	Asn	Va I 210	Pro	Pro	Thr	lle	Arg 215		Arg	GIn	Asn	11e 220		Asn	Ala	Thr
15	Al a 225		Leu	Gly	GIn	Ser 230		Thr	Leu	Val	Cys 235	Asp	Ala	Glu	Arg	Phe 240
20	Pro	Glu	Pro	Thr	Net 245	Ser	Trp	Thr	Lys	Asp 250	Gly	Glu	GIn	lle	@1u 255	Gln
25	Glu	Glu	Asp	Asp 260	Glu	Lys	Tyr	lle	Phe 265	Ser	Asp	Asp	Ser	Ser 270	GIn	Leu
	Thr	He	Lys 275	Lys	Vai	Asp	Lys	Asn 280	Asp	Glu	Ala	Glu	Tyr 285	He	Cys	lle
30	Ala	Glu 290	Asn	Lys	Ala	Gly	Glu 295	Gln	Asp	Ala	Thr	11e 300	His	Leu	Lys	Val
35	Phe 305	Ala	Lys	Pro	Lys	11e 310	Thr	Tyr	Val	Glu	Asn 315	Gin	Thr	Ala	Met	GIu 320
40	Leu	Glu	Glu	Gin	Va I 325	Thr	Leu	Thr		G1u 330	Ala	Ser	Gly		Pro 335	He
45	Pro	Ser	lle	Thr 340	Trp	Arg	Thr		Thr 345	Arg	Asn	lle		Ser 350	Glu	Glu
	Lys	Thr	Leu	Asp	Gly	His	Met	Val	Vaİ	Arg	Ser	His	Ala	Arg '	Val	Ser

5	Ser	7 Lei 37(	u Thi O	r Leu	ı Lys	s Ser	- 11e 37t		Tyr	Thr	- Asp	380		r Glu	ı Tyr	lle
	Cys 385	s Thi	r Ala	a Ser	· Asr	390		∋ Gly	GIn	Asp	Ser 395		s Ser	· Met	Tyr	- Leu 400
10	Glu	ı Val	l Gir	Tyr	Ala 405	Pro	) Lys	s Leu	GIn	Gly 410		Val	Ala	Val	Tyr . 415	Thr
15	Trp	Glu	ı Gly	420	GIn	Val	Asn	ı (le	Thr 425		Glu	Val	Phe	Ala 430		Pro
20 <sub>.</sub>	Ser	Ala	Thr 435		Ser	Trp	Phe	Arg 440	Asp	Gly	GIn	Leu	Leu 445		Ser	Ser
25	Asn	Tyr 450	Ser	Asn	lie	Lys	l le 455		Asn	Thr	Pro	Ser 460		Ser	Tyr	Leu
	G1u 465	Val	Thr	Pro	Asp	Ser 470	Glu	Asn	Asp	Phe	Gly 475	Asn	Tyr	Asn	Cys	Thr 480
30	Ala	Val	Asn	Arg	lle 485	Gly	Gin	Glu	Ser	Phe 490	Glu	Phe	lle	Leu	Va i 495	Gin
35	Ala	Asp	Thr	Pro 500	Ser	Ser	Pro	Ser	lle 505	Asp	GIn	Val	Glu	Pro 510	Tyr	Ser
10	Ser	Thr	Ala 515	GIn	Val	Gln	Phe	Asp 520	Glu	Pro	Glu	Ala	Thr 525	Gly	Gly	Val
15	Pro	lle 530	Leu	Lys	Tyr	Lys	Ala 535	Glu	Trp	Arg	Ala	Va I 540	Gly	Glu	Glu	Val
	Trp 545	His	Ser	Lys		Tyr 550	Asp	Ala	Lys	Glu	Ala 555	Ser	Met	Glu	Gly	11e

5	Va I	Thr	- 116	e Val	Gly 565		Lys	Pro	Glu	Thr 570		Tyr	Ala	∖ Val	Arg 575	Leu
٠.,	Ala	Ala	ı Leu	. Asr 580		Lys	Gly	Leu	Gly 585		He	Ser	Ala	Ala 590		Glu
10	Phe	Lys	Thr 595		) Pro	Val	Gin	Gly 600		Pro	Ser	Ala	Pro 605		Leu	Glu
15	Gly	GIn 610		: Gly	Glu	Asp	Gly 615		Ser	He	Lys	Va I 620		Leu	He	Lys
20	GIn 625	Asp	Asp	Gly	Gly	Ser 630		He	Arg	His	Tyr 635	Leu	Val	Arg	Tyr	Arg 640
25	Ala	Leu	Ser	Ser	Glu 645	Trp	Lys	Pro	Glu	lle 650	Arg	Leu	Pro	Ser	Gly 655	Ser
	Asp	His	Val	Met 660		Lys	Ser	Leu	Asp 665		Asn	Ala	Glu	Tyr 670	Glu	<b>V</b> ạ I
30.	Tyr	Val	Va I 675		Glu	Asn	GIn	GIn 680	Gly	Lys	Ser	Lys	Ala 685	Ala	His	Phe
<b>35</b>	Val	Phe 690	Arg	Thr	Ser	Ala	GIn 695	Pro	Thr	Ala	He	Pro 700	Ala	Asn	Gly	Ser
40	Pro 705	Thr	Ser	Gly	Leu	Ser 710	Thr	Gly	Ala	lie	Val 715	Gly	ile	Leu	He	Va I 720
45	He	Phe	Val	Leu	Leu 725	Leu	Val	Val	Val	Asp 730	He	Thr	Cys	Tyr	Phe 735	Leu
	Asn	Lys	Cys	Gly 740	Leu	Phe	Met		lle 745	Ala	Val	Asn	Leu	Cys 750	Gly	Lys

5	Ala Gly Pro Gly Ala Lys Gly Lys Asp Met Glu Glu Gly Lys Ala Ala 755 760 765	
	Phe Ser Lys Asp Glu Ser Lys Glu Pro IIe Val Glu Val Arg Thr Glu 770 775 780	
10		
	Glu Glu Arg Thr Pro Asn His Asp Gly Gly Lys His Thr Glu Pro Asn 785 790 795 800	
15	Glu Thr Thr Pro Leu Thr Glu Pro Glu Lys Gly Pro Val Glu Ala Lys 805 810 815	
20	Pro Glu Cys Gin Glu Thr Glu Thr Lys Pro Ala Pro Ala Glu Val Lys 820 825 830	
25	Thr Val Pro Asn Asp Ala Thr Gin Thr Lys Giu Asn Giu Ser Lys Ala 835 840 845	
	<210> 9	
	<b>&lt;211&gt;</b> 1692	
	<212> DNA	
30	<pre>&lt;213&gt; Homo sapiens;</pre>	
	⟨220⟩	
	<221> CDS	
	<b>&lt;222&gt;</b> (121) (1080)	
35	<223>	
	<300>	
	<300>	
	<309> 1991-03-19	
10	<313> (1) (1692)	
	<b>&lt;400&gt; 9</b>	
	attcagactg ccagcacttt gctatctaca gccggggctc ccgagcggca gaaagttccg	60
15	gocactotot googottggg ttgggcgaaa gocaggaccg tgccgcgcca ccgccaggat	120
	atg gag cta ctg tcg cca ccg ctc cgc gac gta gac ctg acg gcc ccc	168

	1				5					10					15		
5																gac Asp	216
																ccg Pro	264
10																cac His	312
15																cat His 80	360
20	gtg Val	cgc Arg	gcg Ala	Pro	agc Ser 85	ggg Gly	cac His	cac His	cag Gin	gcg Ala 90	ggc Gly	cgc Arg	tgc Cys	cta Leu	ctg Leu 95	tgg Trp	408
25					Cys											aag Lys	456
30				Met			cgg Arg										504
							tgc Cys 135										552
35							cgc Arg										600
40	cag Gin	gct Ala	ctg Leu	ctg Leu	cgc Arg 165	gac Asp	cag Gin	gac Asp	gcc Ala	gcg Ala 170	ccc Pro	cct Pro	ggc Gly	gca Ala	gcc Ala 175	gcc Ala	648
45	ttc Phe	tat Tyr	gcg Ala	ccg Pro 180	ggc Gly	ccg Pro	ctg Leu	ccc Pro	ccg Pro 185	ggc Gly	cgc Arg	ggc Gly	ggc Gly	gag Glu 190	cac His	tac Tyr	696
							tcc Ser										744

			190					200	1				205				
5	atg Met	atg Met 210	Asp	tac Tyr	agc Ser	ggc	ccc Pro 215	Pro	agc Ser	ggc Gly	gcc Ala	cgg Arg 220	cgg Arg	cgg Arg	aac Asn	tgc Cys	792
10	tac Tyr 225	gaa Glu	ggc Gly	gcc Ala	tac Tyr	tac Tyr 230	aac Asn	gag Glu	gcg Ala	ccc Pro	agc Ser 235	gaa Glu	ccc Pro	agg Arg	ccc Pro	ggg Gly 240	840
10	aag Lys	agt Ser	gcg Ala	gcg Ala	gtg Val 245	tcg Ser	agc Ser	cta Leu	gac Asp	tac Tyr 250	ctg Leu	tcc Ser	agc Ser	atc   le	gtg Val 255	gag Glu	888
15	cgc Arg	atc lle	tcc Ser	acc Thr 260	gag Glu	agc Ser	cct Pro	gcg Ala	gcg Ala 265	ccc Pro	gcc Ala	ctc Leu	ctg Leu	ctg Leu 270	gcg Ala	gac Asp	936
20	gtg Val	cct Pro	tct Ser 275	gag Glu	tcg Ser	cct Pro	ccg Pro	cgc Arg 280	agg Arg	caa Gin	gag Glu	gct Ala	gcc Ala 285	gcc Ala	occ Pro	agc Ser	984
25	gag Glu							Pro									1032
	cag Gln (				Gly					Pro						tga	1080
30	ggggi	gatg	tg g	ccgc	ccaa	c cc	cgcc	aggg	atg	gtgc	cct	aggg	tccc	tc g	cgcc	caaaa	1140
	gatt	gaac	tt a	aatg	cccc	c ct	ccca	acag	cgc	ttta	aaa	gcgc	catc	tc t	tgag	gtagg	1200
35	agagg	gcgg	ag a	actg	aagt	t tc	cgcc	cccc	ccg	acag	ggc	aagg	acac	ag c	gcgg	tttt	1260
	tccad	cgca	gc a	ccct	tctc	g ga	gacc	catt	gcg	atgg	ccg	ctcc	gtgt	tc c	togg	tgggc	1320
40	cagag	gctga	aa c	cttg	aggg	g ct	aggt	tcac	gtt	tctc	gcg	ccct	ccat	gg t	gaga	cctc	1380
	gcaga	accta	aa c	cctg	CCCC	g gg	atgo	accg	gtt	attt	ggg	gggg	cgtg	ag a	cagt	gcact	1440
	ccggt	CCC	aa a <sup>.</sup>	tgta	gcag	g tg	taac	cgta	acc	cacc	occ a	Bacc	cgtti	to c	cggt1	tcagg	1500
15	accac	ttt	tt g	taat	actt <sup>.</sup>	t tt	gtaa <sup>.</sup>	tcta	ttc	ctgt	aaa 1	taaga	agtto	eg ti	ttgc	agag	1560

aggagcccct ggggctgtat ttatctctga ggcagggtgt gtggtgctac agggaatttg

	ta	cgtt	tata	ccg	caggo	ogg (	gcgag	CCE	og ge	gogo	togo	t ca	ggtga	tca	aaa	taaaggo	1680
·	gc	taat	ttat	aa													1692
5	<2 <2	10> 11> 12> 13>	10 319 PRT Homo	o sai	oiens	<b>;</b> ;		-									
10	<b>&lt;4</b>	0 <b>0&gt;</b>	10								•						
15	Me <sup>.</sup>	t Gli	ı Lei	ı Let	Ser 5	· Pro	Pro	Leu	Arg	10	Va!	Asp	Leu	Thr	· Ala 15	Pro	
	Ası	Gly	/ Ser	Leu 20	ı Cys	Ser	Phe	Ala	Thr 25	Thr	Asp	Asp	Phe	Tyr 30	Asp	Asp	
20	Pro	) Cys	Phe 35	Asp	Ser	Pro	Asp	Leu 40	Arg	: Phe	Phe	Glu	Asp 45	Leu	Asp	Pro	
25	Arg	50	Met	His	Va I	Gly	Ala 55	Leu	Leu	Lys	Pro	Glu 60	Glu	His	Ser	His	
30	Phe 65	Pro	Ala	Ala	Val	His 70	Pro	Ala	Pro	Gly	Ala 75	Arg	Glu	Asp	Glu	His 80	
35	Val	Arg	Ala	Pro	Ser 85	Gly	His	His	Gln	A1a 90	Gly	Arg	Cys	Leu	Leu 95	Trp	
	Ala	Cys	Lys	Ala 100	Cys	Lys	Arg	Lys	Thr 105	Thr	Asn	Ala	Asp	Arg 110	Arg	Lys	
40	Ala	Ala	Thr 115	Met	Arg	Glu	Arg	Arg 120	Arg	Leu	Ser	Lys	Va I 125	Asn	Glu	Ala	
45	Phe	Glu 130	Thr	Leu	Lys	Arg	Cys 135	Thr	Ser	Ser	Asn	Pro 140	Asn	GIn	Arg	Leu	

	145	-yo v	21 (1)	u iie	150	i Ari	g Asi	) Ala	3     (	3 Arg 155		·    (	e Glu	ı Giy	160
5	Gin A	lla Le	eu Lei	u Arg 165	Asp	Gir	ı Asp	Ala	170		Pro	Gly	/ Ala	Ala 175	
10	Phe T	yr Ai	a Pro 180	o Gly	Pro	Leu	ı Pro	Pro 185	Gly	' Arg	: Gly	Gly	Glu 190		Tyr
15	Ser G	ly As 19	p Ser 5	Asp	Ala	Ser	Ser 200	Pro	Arg	; Ser	Asn	Cys 205		Asp	Gly
	Met Met 2	et As 10	p Tyr	Ser	Gly	Pro 215	Pro	· Ser	Gly	Ala	Arg 220	Arg	Arg	Asn	Cys
20	Tyr Gi 225	lu Gl	y Ala	Tyr	Tyr 230	Asn	Glu	Ala	Pro	Ser 235	Glu	Pro	Arg	Pro	Gly 240
25	Lys Se	er Ala	a Ala	Va I 245	Ser	Ser	Leu	Asp	Tyr 250	Leu	Ser	Ser	ile	Va I 255	Glu
30	Arg	e Ser	Thr 260	Glu	Ser	Pro	Ala	Ala 265	Pro	Ala	Leu	Leu	Leu 270	Ala	Asp
35	Val Pr	o Ser 275	Glu	Ser	Pro	Pro	Arg 280	Arg	Gln	Glu		Ala 285	Ala	Pro :	Ser
	Glu Gl	y Glu D	Ser	Ser	Gly /	Asp 295	Pro	Thr	GIn		Pro . 300	Asp	Ala	Alai	Pro
40.	GIn Cys	s Pro	Ala	Gly /	Ala / 310	Asn	Pro .	Asn		lle ' 315	Tyr (	Gin	Val I	Leu	
15		11 1427 DNA Homo	sapi	ens;											,

. 5	<2 <2	20> 21> 22> 23>	CDS (43	)(	810)												
10	<30 <30	00> 08> 09> 13>	200	00559 3-04- (14	<b>-</b> 07												
	<40	00>	11														
	CC1	tctc	gctg	ccg	tcca	ggt g	caco	gcc	tg co	ctctc	cagca	gg			gtg		54
15													Met 1	Asp	Val	Met	
20	gat Asp 5	ggc Gly	tgo Cys	cag Gir	tto Phe	tca Ser 10	cct Pro	tct Ser	t gag · Glu	tac I Tyr	tto Phe	tac Tyr	gad Asp	gge Gly	tcc Ser	tgc Cys 20	102
	ata Ile	o CCg Pro	tco Ser	Pro	gag Glu 25	ggt Gly	gaa Glu	ttt Phe	ggg Gly	gac Asp 30	gag Glu	ttt Phe	gtg Val	ccg Pro	cga Arg 35	gtg Val	150
25	gct Ala	gcc Ala	ttc Phe	gga Gly 40	gcg	cac His	aaa Lys	gca Ala	gag Glu 45	ctg Leu	cag Gin	ggc Gly	tca Ser	gat Asp 50	gag Glu	gac Asp	198
30	gag Glu	cac His	gtg Val 55	cga Arg	gcg Ala	cct Pro	acc Thr	ggc Gly 60	cac His	cac His	cag Gin	gct Ala	ggt Gly 65	cac His	tgc Cys	ctc Leu	246
35	atg Met	tgg Trp 70	gcc Ala	tgc Cys	aaa Lys	gcc Ala	tgc Cys 75	aag Lys	agg Arg	aag Lys	tcc Ser	acc Thr 80	acc Thr	atg Met	gat Asp	cgg Arg	294
40	cgg Arg 85	aag Lys	gca Ala	gcc Ala	act Thr	atg Met 90	cgc Arg	gag Glu	cgg Arg	agg Arg	cgc Arg 95	ctg Leu	aag Lys	aag Lys	gtc Val	aac Asn 100	342
<b>-</b> ‡ U	cag Gin	gct Ala	ttc Phe	gaa Glu	acc Thr 105	ctc Leu	aag Lys	agg Arg	tgt Cys	acc Thr 110	acg Thr	acc Thr	aac Asn	ccc Pro	aac Asn 115	cag GIn	390
45	agg Arg	ctg Leu	Pro	aag Lys 120	gtg Val	gag Glu	atc lle	Leu	agg Arg 125	aat Asn	gcc Ala	atc Ile	Arg	tac Tyr 130	atc   e	gag Glu	438

	agc ctg Ser Lei	cag Gin 135	gag Glu	ttg Leu	ctg Leu	aga Arg	gag Glu 140	Gin	gtg Val	gag Glu	aac Asn	tac Tyr 145	tat Tyr	agc Ser	ctg Leu	486
5	ccg ggg Pro Gly 150	Gln	agc Ser	tgc Cys	tcg Ser	gag Glu 155	ccc Pro	acc Thr	agc Ser	coc Pro	acc Thr 160	tcc Ser	aac Asn	tgc Cys	tct Ser	534
10	gat ggd Asp Gly 165	atg Met	ccc Pro	gaa Glu	tgt Cys 170	aac Asn	agt Ser	cct Pro	gtc Val	tgg Trp 175	tcc Ser	aga Arg	aag Lys	agc Ser	agt Ser 180	582
15	act ttt Thr Phe	gac Asp	Ser	atc   e  185	tac Tyr	tgt Cys	cct Pro	gat Asp	gta Val 190	tca Ser	aat Asn	gta Val	tat Tyr	gcc Ala 195	aca Thr	630
20	gat aaa Asp Lys	Asn														678
20	cgg atc Arg lle					GIn					Leu					726
25	tct ctc Ser Leu 230				Ala											774
30	gct tct Ala Ser 245			Arg I					Val		tga	acta	attt	tc		820
,	tggtctat	at ga	actto	cttc	c ag	gagg	gcct	aat	acac	agg	acga	agaa	gg c	ttca	aaaag	880
35	toccaaac	ca a	gacaa	acat	g tạ	cața	aaga	ttt	cttt	tca	gttg	taaa	tt t	gtaa	agatt	940
	accttgcc	ac ti	ttata	aagaa	a ag	tgta <sup>.</sup>	ttta	act	aaaa	agt	catc	attg	ca a	ataa	tactt	1000
40	tottotto	tt ta	attat	ttct1	t tgo	otta	gata	tta	atac	ata :	gttc	cagt	aa ta	acta	tttct	1060
	gatagggg	gc ca	attga	attge	a ggg	gtago	ottg	ttc	gaat	gct	taact	ttat	at at	tacat	tatat	1120
	atatatta	ta aa	atatt	goto	ato	2888	atgt	ctc	tggt	gtt ·	tagag	gotti	ta ti	tttt	ttctt	1180
45	taaaacat	ta aa	acag	ctga	gae	atcag	tta	aat	ggaa <sup>.</sup>	ttt 1	taaat	tata	tt te	acta	attto	1240
	ttttctct	tt aa	itcct	ttag	tte	itati	:gta	ttaa	ata	aa a	atata	atad	et go	ctae	tgta	1300

	tatattttga	tottttottg	taagaaatg	t atcttttaaa	tgtaagcaca	aaatagtact 1360
	ttgtggatca	tttcaagata	taagaaatt	t tggaaattcc	accataaata	aaatttttta 1420
5	ctacaag					1427
	<210> 12					
10	<211> 255 <212> PRT					
10		o sapiens:				
	<b>&lt;400&gt;</b> 12					
15	Met Asp Va 1	l Met Asp G 5	ly Cys Gin	Phe Ser Pro S	Ser Glu Tyr	Phe Tyr 15
20	Asp Gly Se	r Cys lle Pi 20	o Ser Pro	Glu Gly Glu I 25	Phe Gly Asp 30	Glu Phe
25	Val Pro Arg 35	g Val Ala Al	a Phe Gly 40	Ala His Lys A	Ala Glu Leu 45	Gin Gly
	Ser Asp Glu 50	ı Asp Glu Hi	s Val Arg 55	Ala Pro Thr 6	aly His His 60	Gin Ala .
30	Gly His Cys 65	S Leu Met Tr 70	p Ala Cys	Lys Ala Cys L 75	ys Arg Lys	Ser Thr 80
35	Thr Met Asp	Arg Arg Ly 85	s Ala Ala	Thr Met Arg G 90		Arg Leu 95
40 .	Lys Lys Val	Asn Gin Ala		Thr Leu Lys A 105	rg Cys Thr 110	Thr Thr
45	Asn Pro Asn 115	Gin Arg Let	ı Pro Lys \ 120	Val Glu lle Lo	eu Arg Asn <i>l</i> 125	Ala lle
	Arg Tyr ile 130	Glu Ser Leu	Gin Glu L 135	eu Leu Arg 6.	lu Gin Vai 6 40	ilu Asn

140

5	1yr 145	Tyr	- Ser	Leu	1 Pro	150	Gin	Ser	Cys	Ser	155		Thr	Ser	Pro	160
	Ser	' Asn	Cys	Ser	Asp 165	Gly	Met	Pro	Glu	Cys 170		Ser	Pro	Val	Trp 175	Ser
10	Arg	Lys	: Ser	Ser 180	Thr	Phe	Asp	Ser	l le 185	Tyr	Cys	Pro	Asp	Val 190		Asn
15	Val	Tyr	Ala 195	Thr	Asp	Lys	Asn	Ser 200	Leu	Ser	Ser	Leu	Asp 205	Cys	Leu	Ser
20	Asn	lle 210	Val	Asp	Arg	lle	Thr 215	Ser	Ser	Glu	GIn	Pro 220	Gly	Leu	Pro	Leu
25	GIn 225	Asp	Leu	Ala	Ser	Leu 230	Ser	Pro	Vai	Ala	Ser 235	Thr	Asp	Ser	Gin	Pro 240
	Arg	Thr	Pro	Gly	Ala 245	Ser	Ser	Ser	Arg	Leu 250	He	Tyr	His	Val	Leu 255	
30	4040		_													
	<b>&lt;210</b>	_	3													
	<211 <212		175 Na													
	<212 <213			sapi	ono'				•							
35	\0	, ,		зарт	G110,											
	<220	>														
	<b>&lt;221</b>	> c	DS													
	<b>&lt;222</b>	> (	1)	(675)	)											
	<b>&lt;223</b>	>														
10	4															
	<b>&lt;300</b>															
	<308)		T007:													
	<309) <313)	_		05-13 (675)												
5	\U 1 U/	• (	./	(0/0/	•	•										
-	<b>&lt;400</b>	> 1:	3													
	atg g			tat g	ag e	ica 1	toc a	300 1	tac ·	tte 1	tac 4	iae i	788	200	cac .	tta
	Met (	alu I	_eu ]	Tyr G	lu 1	Thr S	Ser F	ro 1	ſyr I	Phe	Tyr (	3In (	3lu i	Pro A	Arg	Phe

	ı				5					10					15		
5	ta Ty	t ga r As	nt gg p Gi	g ga y Gi 20	aA u	o ta n Ty	c ct; r Lei	g cc u Pr	t gto o Va 25	c cad	c ct s Le	c ca u Gli	g gg n Gly	c tt y Ph 30	e GI	a cca u Pro	96
10	eca Pro	a gg o Gi	c ta y Ty 35	r GI	g cg u Ar	g ac <sub>i</sub>	g gag r Glu	cto Leo 40	c acc u Thr	cti Lei	g ago	c ccc r Pro	c gag o Glu 45	g gc I Al	c cc a Pr	a ggg o Gly	. 144
	Pro	ct Le 50	u GI	g ga u As <sub>i</sub>	c aa p Ly	g ggg s Gly	ctg Leu 55	ggi Gly	g acc	e ccc	gaş Glu	g cad His 60	tgt Cys	: cc	a ggo o Gly	c cag y Gin	192
15	tgo Cys 65	ct: Lei	g co u Pro	g tgg o Trg	g go; o Ala	g tgl a Cys 70	: aag : Lys	gte Val	g tgt I Cys	: aag : Lys	agg Arg 75	g aag g Lys	tcg Ser	gtį Val	g too I Ser	gtg Val 80	240
20	gac Asp	cgg Arg	g cgg	g ogg g Arg	g gcg Ala 85	g goo	aca Thr	ctg Leu	g agg I Arg	gag Glu 90	: aag Lys	cgc Arg	agg Arg	Leu	aag Lys 95	aag Lys	288
25	gtg Val	aat Asr	gag Glu	g gco 1 Ala 100	Phe	gag Glu	gcc Ala	ctg Leu	aag Lys 105	Arg	agc Ser	acc Thr	ctg Leu	cto Leu 110	Asn	ccc Pro	336
30	aac Asn	cag Gin	cgg Arg 115	Leu	Pro	aag Lys	gtg Val	gag Glu 120	He	ctg Leu	cgc Arg	agt Ser	gcc Ala 125	ato	cag Gin	tac Tyr	384
	atc Ile	gag Glu 130	Arg	ctc Leu	cag Gin	gcc Ala	ctg Leu 135	ctc Leu	agc Ser	tcc Ser	ctc Leu	aac Asn 140	GIn	gag Glu	gag Glu	cgt Arg	432
35	gac Asp 145	ctc Leu	cgc Arg	tac Tyr	cgg Arg	ggc Gly 150	ggg Gly	ggc Gly	ggg Gly	ccc Pro	cag GIn 155	cca Pro	ggg Gly	gtg Val	ccc Pro	agc Ser 160	480
40	gaa Glu	tgc Cys	agc Ser	tct Ser	cac His 165	agc Ser	gcc Ala	tcc Ser	tgc Cys	agt Ser 170	cca Pro	gag Giu	tgg Trp	ggc Gly	agt Ser 175	gca Ala	528
<b>45</b> .	ctg Leu	gag Glu	ttc Phe	agc Ser 180	gcc Ala	aac Asn	cca Pro	ggg Gly	gat Asp 185	cat His	ctg Leu	ctc Leu	Thr	gct Ala 190	gac Asp	cct Pro	576
	aca . Thr .	gat Asp	gcc Ala	cac His	aac Asn	ctg Leu	cac His	tcc Ser	cto Leu	acc Thr	tcc Ser	atc Ile	gtg Val	gac Asp	agc Ser	atc 11e	624

672

675

195 200 205

aca gtg gaa gat gtg tct gtg gcc ttc cca gat gaa acc atg ccc aac Thr Val Glu Asp Val Ser Val Ala Phe Pro Asp Glu Thr Met Pro Asn 210 220

tag

10 <210> 14 <211> 224 <212> PRT <213> Homo sapiens;

15 <400> 14

40

Met Glu Leu Tyr Glu Thr Ser Pro Tyr Phe Tyr Gln Glu Pro Arg Phe 1 5 10 15

Tyr Asp Gly Glu Asn Tyr Leu Pro Val His Leu Gin Gly Phe Glu Pro
20 25 30

- Pro Gly Tyr Glu Arg Thr Glu Leu Thr Leu Ser Pro Glu Ala Pro Gly 35 40 45
- Pro Leu Glu Asp Lys Gly Leu Gly Thr Pro Glu His Cys Pro Gly Gln 30 50 55 60
- Cys Leu Pro Trp Ala Cys Lys Val Cys Lys Arg Lys Ser Val Ser Val 65 70 75 80

Asp Arg Arg Ala Ala Thr Leu Arg Glu Lys Arg Arg Leu Lys Lys 85 90 95

- Val Asn Glu Ala Phe Glu Ala Leu Lys Arg Ser Thr Leu Leu Asn Pro 100 105 110
- 45 Asn Gin Arg Leu Pro Lys Val Giu IIe Leu Arg Ser Ala IIe Gin Tyr 115 120 125

	lle	Glu 130	Arg	Leu	Gin	Ala	Leu 135	Leu	Šer	Ser	Leu	Asn 140	Gin	Glu	Glu	Arg	
5	Asp   145 <sup>-</sup>	Leu	Arg	Tyr	Arg	Gly 150	Gly	Gly	Gly	Pro	Gin 155	Pro	Gly	Vai	Pro	Ser 160	
10	Glu (	Cys	Ser	Ser	His 165	Ser	Ala	Ser	Cys	Ser 170	Pro	Glu	Trp	Gly	Ser 175	Ala	
15	Leu (	Glu	Phe	Ser 180	Ala	Asn	Pro	Gly	Asp 185	His	Leu	Leu	Thr	Ala 190	Asp	Pro	
	Thr /	Asp	Ala 195	His	Asn	Leu	His	Ser 200	Leu	Thr	Ser	lie	Va I 205	Asp	Ser	lle	
20	Thr V	/a l 210	Glu	Asp	Val	Ser	Val 215	Ala	Phe	Pro		Glu 220	Thr	Met	Pro	Asn	
<b>2</b> 5	<210> <211> <212> <213>	2 D	6 Na	icia	ıl Se	quen	се										
30	<220> <223>		RY fo	orwa	rd p	rime	r										
35	<400> gcctc	• • •	-	tatt	aatc	t ct	ggag										26
10	<210> <211> <212> <213>	23 DN	B VA	icia	l Sec	quen	C <del>0</del>										
	<220> <223>	SR	lY r∈	over	se pi	rime	r										
15	<400> gctgat			latto	otgoa	a tgo	3							٠			23

```
<210> 17
      <211> 26
      <212> DNA
      <213> Artificial Sequence
  5
      <220>
      <223> SRY probe
      <400> 17
 10
      aggcgcaagt tggctcaaca gaatcc
                                                                          26
      ₹210> 18
      <211> 25
      <212> DNA
15
      <213> Artificial Sequence
      <220>
      <223> IL2 forward primer
20
      <400> 18
      gccttgtgtg ttataagtag gaggc
                                                                          25
25
      <210> 19
      <211> 21
      <212> DNA
     <213> Artificial Sequence
     <220>
30
     <223> IL2 reverse primer
     <400> 19
     agtgccaatt cgatgatgag c
                                                                         21
35
     <210> 20
     <211> 29
     <212> DNA
     <213> Artificial Sequence
40
     <220>
     <223> IL2 probe
     <400> 20
45
     tctcctcaga aattccacca cagttgctg
                                                                         29
```